

No.	Score	Match	Length	DB	ID	Description
1	751.6	24.1	1857	3	US-09-332-478-24	Sequence 24, App
2	751.6	24.1	1857	4	US-09-586-106D-24	Sequence 24, App
3	90	2.9	7218	1	US-08-232-463-14	Sequence 14, App
4	63.8	2.0	1166	3	US-09-072-596-323	Sequence 323, App
5	63.8	2.0	1166	4	US-09-072-967-328	Sequence 328, App
6	63.8	2.0	7218	1	US-08-232-463-14	Sequence 14, App
7	59.8	1.9	37155	4	US-09-949-016-16945	Sequence 16945, App
8	55	1.8	53526	3	US-08-658-136-2	Sequence 2, App
9	55	1.8	53577	3	US-08-658-136-1	Sequence 1, App
C 10	51.4	1.6	767677	4	US-09-949-016-12147	Sequence 12147, App
C 11	51.4	1.6	767677	4	US-09-949-016-17351	Sequence 17351, App
C 12	50.6	1.6	18955	4	US-09-949-016-15780	Sequence 15780, App
C 13	50.6	1.6	30678	4	US-09-949-016-12818	Sequence 12818, App
14	49.8	1.6	234884	4	US-09-949-016-16420	Sequence 16420, App
15	48.6	1.6	86414	4	US-09-949-016-12345	Sequence 12345, App
16	48.6	1.6	86414	4	US-09-949-016-15758	Sequence 15758, App
17	47.4	1.5	390894	4	US-09-949-016-14730	Sequence 14730, App
18	47	1.5	601	4	US-09-949-016-135107	Sequence 135107, App
19	47	1.5	117807	4	US-09-949-016-15555	Sequence 15555, App
20	47	1.5	110332	4	US-09-949-016-12160	Sequence 12160, App
21	47	1.5	115032	4	US-09-949-016-17268	Sequence 17268, App
22	46.8	1.5	247781	4	US-09-949-016-14133	Sequence 14133, App
23	45.4	1.5	67002	4	US-09-949-016-16803	Sequence 16803, App
24	45	1.4	9293	4	US-09-949-016-16801	Sequence 16801, App
C 25	44.4	1.4	81384	4	US-09-949-016-12900	Sequence 12900, App
C 26	44	1.4	46589	4	US-09-949-016-12942	Sequence 12942, App
27	44	1.4	476044	4	US-09-949-016-12412	Sequence 12412, App

C 28	43.6	1.4	19438	4	US-09-949-016-12699	A	Sequence 12699, A
C 29	43.6	1.4	46823	4	US-09-949-016-12723	A	Sequence 12723, A
C 30	43.6	1.4	46940	4	US-09-949-016-16552	A	Sequence 16552, A
C 31	42.8	1.4	15292	4	US-09-949-016-13584	A	Sequence 13584, A
C 32	42.6	1.4	318	3	US-09-165-264-12	App	Sequence 12, App
C 33	42.8	1.4	27687	4	US-09-949-016-13340	A	Sequence 13340, A
C 34	41.8	1.3	68580	4	US-09-949-016-15444	A	Sequence 15840, A
C 35	41.6	1.3	32207	2	US-08-1770-379-20	App	Sequence 20, App
C 36	41.6	1.3	32207	3	US-08-1757-669A-20	App	Sequence 20, App
C 37	41.6	1.3	32207	3	US-09-2330-371A-20	App	Sequence 20, App
C 38	41.6	1.3	16934	4	US-09-949-016-13720	A	Sequence 13720, A
C 39	41.4	1.3	114139	4	US-09-949-016-15536	A	Sequence 16536, A
C 40	41	1.3	42053	4	US-09-949-016-15264	A	Sequence 15924, A
C 41	41	1.3	85675	4	US-09-949-016-13333	A	Sequence 12333, A
C 42	41	1.3	85675	4	US-09-949-016-15855	A	Sequence 15956, A
C 43	40.8	1.3	43414	4	US-09-949-016-12839	A	Sequence 12839, A
C 44	40.8	1.3	43414	4	US-09-949-016-16491	A	Sequence 16491, A
C 45	40.8	1.3	109610	4	US-09-949-016-13525	A	Sequence 13525, A

ALIGNMENTS

RESULT 1
 US-09-322-478-24
 Sequence 24, Application US/09322478
 Patent No. 631662
 GENERAL INFORMATION:
 APPLICANT: Wright, David A.
 APPLICANT: Voytas, Daniel F.
 TITLE OR INVENTION: Plant Retroelements and Met
 FILE REFERENCE: P-1065 ISRF Plant Retroelement
 CURRENT APPLICATION NUMBER: US/09/322,478
 CURRENT FILING DATE: 1999-05-28
 EARLIER APPLICATION NUMBER: 60/087125
 EARLIER FILING DATE: 1998-05-29
 NUMBER OF SEQ ID NOS: 41
 SOFTWARE: PatentIn Ver. 2.0
 SEQ ID NO 24
 LENGTH: 1857
 TYPE: DNA
 ORGANISM: Arabidopsis thaliana
 US-09-322-478-24

Query Match	24.1%;	Score 751.6;	DB 3;	Length 1857;
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Matches 1250; Conservative 0; Mismatches 564; Indels 100; Gaps 6;

Qy	898	ATAGTAACTACAGATGGAAAAATCCTCATATGAGACCCCTATTAATATGTGGATGAAGCTAAG	957
Db	1	ATAGCAATTAACAGATGGCACTTCTTCTGTGATCTTGACTACATCAATATGATGAAGACGAA	60
Qy	958	TCCTGGTCCAATGACCGGAGTGGAGCAACATTTTACGAGAGCTTATAGGATGAATTT	1017
Db	61	TCGTATCTTCAGAGCCAGAGAGAAACAGAGAAATACGAAAGTTTCAG-----	110
Qy	1018	GAACGCTCTGACAGCTGACGTAAATCAAAAGAGAGCTGAAATCGGTAGAGGAAAGGCG	1077
Db	111	-----AAGGAAAGCTGAGATAGCCCGAGGAAAGAGCG	144
Qy	1078	ATGTCCAGTGAATATGAGCTGATTTGATGAGAGATATCAAAACTAGTATGAGCCAGATCA	1137
Db	145	ATGAGAGAGAGTATGAGCTTATATGACGAAGATCTGGAGGACGAGTACATCTCTGAACAG	204
Qy	1138	TGGCGCAAGAGACGAAGCTACTGAACAAATCCGACGAGTTATAGTGGAGAGATATTC	1197
Db	205	ACTGCGAGAGCTACCAACTTCTGCAAGAGCCGACATATTGCTCTGAGGAATATGTT	264
Qy	1198	AGATTCTTTGATGAATGACTTCTGGGGAACGAGTATCCCTGATATGAGACTTTAGCC	1257
Db	265	AGCGTTTTCAAGCTGATGATGAGTTCTGTAGCAGAGGTATCTTCTCGAAGCTCACTTCA	324
Qy	1258	CAGTTGGGGTTACTGAGAGACGTACAGACTTCGTTCCAGAAAGTCACTC-----	1308

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Db      325  CAACTGGAGTGTGGAAAGATGTGACGACCTGTACCAAAAGTTGTCATCTGACACTTTG 384
Qy      1309  -----ATAAGAGAGAGACATGAGTTTCTTTCCACACTGCA 1346
Db      385  ATGGCTATCCGATATGATAGCATATGAAATGAGACAAATACATCTCTCCACATACAA 444
Qy      1347  GTGAAATGTATGAGGAGACTCAACACTTTGAGCTGAGATCAATGGGGTTAGGCTTTG 1406
Db      445  GTAGAGCTCTACCAAGGTATGACCTCTGATGAGTTGATTTGAAAGATTTGGATTTCTG 504
Qy      1407  ACGTCTTAGTGTGAAACAGGGGTACAGATTTTATGATGAAGAATTTGGAAGACGTGT 1466
Db      505  CGATTTCTGTATGATGATAGTACAGGTATCAATCAACGATTTGGAAGATTTGTT 564
Qy      1467  GGTTCCTCCAGTGTGAAAGGAAACCAACCCAGTTTTCAGAGGAAAGCTTAAAGATTG 1526
Db      565  GATTTTCCAGTGTGAAACGGGATCTTAAGCAAAATGTAAAGAAAGATTTGAAAGACTTG 624
Qy      1527  TGGGCTACTATTGGGAAACATCTACCGCTAACTCGACGGGTCCAGAGCAACCAATC 1586
Db      625  TGGATCACCATGTGGAGCTCTGTACCGTTGAATGCTTCAGGGTCAAAAGACATCAGAT 684
Qy      1587  CGGAGTCCCTGTGATTCGCTACTTTGAGCGCTGGTTGCCAATGTTTTTAATCCAGGAG 1646
Db      685  CGACCCCTGTATCAGGACTTCCAGCGTCTGTAGCCACAGTACTTACTCCGAGAG 744
Qy      1647  TCTACAGGACCCGTGTCTTACACAGACATGAAATGATGATTCAGCGCTTAAAGGATT 1706
Db      745  ATTACAGGAGCTGTCACTTACTCTGATATGAGATGATGCAATGGCCCTCAAGAAACT 804
Qy      1707  CTCGCCCTTACAAAAGAAAGATGCTCTGAGAGAGATCTTACGACTCACCACTG 1766
Db      805  CTCGCCCAAACTTAAATGAGCATGTCTCCACAGGGTGAAGTCAATGACACACTCTCT 864
Qy      1767  ATGCTCTGTGATCATCTGTGTGGGTACATGAAGTGGCGCTGCAACAGGCAAG 1826
Db      865  ATACTCTCTGTATCATCTGTGTGATACAAAACCTGGCGGTGCAATTAACGCAAG 924
Qy      1827  AAGGTAAAGAGACACTATCGTGGGTGCGCTTTGTCACCCCAATTCGAAATTTGTGA 1886
Db      925  AGAGACAGAGCGGCTGTGTGATAGGTGCGGTGTGACACCTATCTGTACTGTGTGA 984
Qy      1887  GTTCCGCTCAGAAAGTATGAGGTTAGACACCGAATGATGACTTGGATCTTGGCCGA 1946
Db      985  GTCCCACTCATTTCTGTGACTCGACGCCACAGCATGATGATGAGCACTTACGCTC 1044
Qy      1947  TGTGAGTCTGTGATTTGACATGATGTGGCGACTTTACCCGCTACAGTTTCAGATTTC 2006
Db      1045  TGCCAATTCGTGAGATTTGCAATGTGTGACATTTCCACAGGTTCAAGTTTGAACA 1104
Qy      2007  TCGATTAGATTCGCAACATTTCTTTCCCTGCAATTAAGTACTAGATTCTCGAGG 2066
Db      1105  ACAAGACAGAGAGCTTACATCTTCTCCCTAGCCCTGAGGTACACAGATTAATCAGG 1164
Qy      2067  AGGAACATTGACTTCAAGCTCGCTGTAAGATCTTTATTTTGAAGGAGTCCGCCACT 2126
Db      1165  GATTAATTTGATTTTAAAGCTTGAATTTGACCGCTCTACTATGAGAACCTCCACAT 1224
Qy      2127  GAGGAGATTAGTCAACCGAAGAGCTTCAATTAAGAAATGTTGATGAGACATATGATA 2186
Db      1225  GATGAGAGCATCTTCTGAAAGACT-----GCTTGGATGGAGTGAATGAAGT 1275
Qy      2187  GATGAGGCGAGTTTGAACAGAGCATGTATCAATTTCAAGAGACATTAATCTCAGAG 2246
Db      1276  GGAGAGATTAAGTTTGAACATGATGATCTTCTGCTGCAACATGATCTCAGAGAG 1335
Qy      2247  AAAAGCAAGATTTGAGCGAGCTCAACAGAAACAAGCAAGCTGCAAGTGTGCAAG 2306
Db      1336  CAGACCAAGAGCTTGAAGCTCAATAGAAATTAAGATTAATTTGAGAAAGTGTGCA 1395
Qy      2307  AAACAGATTAAGTACTGCGCAAGTGCCTCAGGGCTATGAATTTCTGAGAGCAAGATC 2366

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Db      1396  AAGCAGACAGGCTGATGCCAAGTG-----TTTCAAGCTTCTGACAGACAGACTG 1446
Qy      2367  AGTGTCTCTCTTCCACTTACAACTATTTCCGAAATGACAGCTCCCTCAGAGACATG 2426
Db      1447  AGTGTCTCTCTCCACACATGCTATTTCCAGAGTAAACCTCTATGAAATGCAATG 1506
Qy      2427  AGGAATATGACG-----GCCGAGCTTAAGAGCAGAAAGATTTGATGTG 2474
Db      1507  AGGAATTAATTAATGACCTGCGCACAGGCTTGAAGCTTACGAGCAGAGATGCCCAT 1566
Qy      2475  CTTGCGAGCAATTCATCATTTGAGGCTGCTGATCTAGAAAGATTAAGAAACGACACT 2534
Db      1567  CAGGCTAGGCAATTTCTCATTTGAAATCCCGGAAACAGAAAGAAAGAGCTTACATC 1626
Qy      2535  ACTGATCTAGACAGACAGAGACAGACTTGTGACGTCTGTAGTTTACGAGCCGAGT 2594
Db      1627  ACTGATCTTAGAGC---AGATCACGGCTCATTTCACTGAGAGATCACTGACCGTGT 1683
Qy      2595  GTTGGCCGCAATTAAGAAAGAGAGGTGAGATTTCTCAGAGCGGTGCGCCACAGA 2654
Db      1684  GTTGGCCGACAGAGAGAGAGATGTGAGATTTCTCAGAGCGGTGCGCCACAGA 1743
Qy      2655  GCTGATGATGATGATGATCCCATGCTGAGCTGATACGAAACATGGCGGTTGCTATG 2714
Db      1744  GCTGATGAGTGTGATGATCCCATGCTGCTGAGCTGATACAGAAACAGAGGTTGCTAT 1803
Qy      2715  GCTTGGAGCAATCAAGAGAGCCCATGACTTACCAACTTGTCTTATTTGAC 2768
Db      1804  GCTTGGAGCAATGACAGAGCCATTAACGAGCAATGATGATTTCTTGAC 1857

RESULT 2
US-09-586-106D-24
; Sequence 24, Application US/09586106D
; Patent No. 6720479
; GENERAL INFORMATION:
; APPLICANT: Wright, David A.
; APPLICANT: Voytas, Daniel F.
; TITLE OF INVENTION: PLANT RETROELEMENTS AND METHODS RELATED THERETO
; FILE REFERENCE: P-1065A
; CURRENT APPLICATION NUMBER: US/09/586,106D
; PRIOR FILING DATE: 2003-02-07
; PRIOR APPLICATION NUMBER: 60/087,125
; PRIOR FILING DATE: 1998-05-29
; PRIOR APPLICATION NUMBER: 09/322,478
; PRIOR FILING DATE: 1999-05-28
; NUMBER OF SEQ ID NOS: 190
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 24
; LENGTH: 1857
; TYPE: DNA
; ORGANISM: Arabidopsis thaliana
US-09-586-106D-24

Query Match      24.1%; Score 751.6; DB 4; Length 1857;
Best Local Similarity 65.3%; Pred. No. 3.4e-217;
Matches 1250; Conservative 0; Mismatches 564; Indels 100; Gaps 6;

```


NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 30472/114 IMMUN
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 836-9300
TELEFAX: (703) 683-4109
TELEX: 899149
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 7218 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
IMMEDIATE SOURCE:
CLONE: pTZapT-Fls
US-08-232-463-14

Query Match 2.9%; Score 90; DB 1; Length 7218;
Best Local Similarity 4.5%; Pred. No. 1.4e-15;
Matches 18; Conservative 253; Mismatches 133; Indels 0; Gaps 0;

50 AAGCCGATCGCTTCCCTCACAACCTCTCACTGACGCGCGCTCTCTCACTTACT 109
1053 AGGAGCTTGGATTT 1112

110 CGGCTTATCGCTCTCATCGCATCTCTCAACATACTGACCTCGGATATCACTGAGC 169
1113 TT 1172

170 TCGCGCTTTCACGCGCTTCACGCTGTCACGCGCGCTCTCTCAAGAAACAAC 229
1173 TT 1232

230 TCGAGCTCTCATCTCTCACTGACCTCTACACCAAGCGGCTTCACACTTCTAGC 289
1233 TT 1292

290 TCTTAAACATCGACCACTTCACATCAACCAATCAATGTTTCTCTCTCAATTAAG 349
1293 TT 1352

350 CTGACATACGACGCGTGAACACTTACACTCAAGCTCTCATCTCTCACTGTTT 409
1353 TT 1412

410 CCAACACGCTGCTCTCATCTCCCAAGCAAGCTTCTCACTGACT 453
1413 TT 1456

RESULT 4
US-09-072-596-323
Sequence 323, Application US/09072596
Patent No. 6458366
GENERAL INFORMATION:
APPLICANT: Reed, Steven G.
APPLICANT: Skelky, Yasir A.W.
APPLICANT: Dillon, David C.
APPLICANT: Campos-Neto, Antonia
APPLICANT: Houghton, Raymond
APPLICANT: Vedvick, Thomas S.
APPLICANT: Twardzik, Daniel R.
APPLICANT: Lodes, Michael J.
APPLICANT: Hendrickson, Ronald C.
TITLE OF INVENTION: COMPOUNDS AND METHODS FOR DIAGNOSIS OF TUBERCULOSIS
NUMBER OF SEQUENCES: 350
CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
CITY: Seattle
STATE: Washington
COUNTRY: USA
ZIP: 98104-7092

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/072,596
FILING DATE: 05-MAY-1998
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Maki, David J.
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 210121.417C9
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
TELEFAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 323:
SEQUENCE CHARACTERISTICS:
LENGTH: 1166 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: Genomic DNA
US-09-072-596-323

Query Match 2.0%; Score 63.8; DB 3; Length 1166;
Best Local Similarity 26.7%; Pred. No. 3.2e-08;
Matches 190; Conservative 136; Mismatches 384; Indels 1; Gaps 1;

2 TCATATATGACGCTTCTCTCATTTCTTGATTCGAAACAAACAAACGCGCATTC 61
373 KSAAMTSMKMGSTSYCTMTYCNNGASTAMTYNNCCCCGMAWCKSMAWCCCTGCA 432

62 CTTCCTCCCAACATCTCACTGACACGCGCGCTCTCACTTACTCGGCTTCATGC 121
433 TYCCMCGSSGXYCTCAGNCCAGCTGNGYCCCTCCMKTCTCACTCACTCGGTTCC 492

122 TCTCATGCGATCTCTCAACATCACTGACCTCGGATATCACTGAGCTGCGCTTCTC 181
493 TTTMMNCCSCNCRCTCAGMCTKSGKACANATMYCCSACKGHTCTMYMSCAKMNT 552

182 ACCGCTTTCATGCTGACGCGCGCTGCTCTCTCTCCCAAGAAACAACATCGAGCTTCCA 241
553 TCCTCTCCTCTTNNCCMCMCSCTTMTMAACTCKCGGAYCKNOMYCTCTCKCAY 612

242 TTTCACTCACTGACCTCTCAACCAACGCGGCTTCACTTACTTAAACATCTC 301
613 NMAACCKTCTYCMGMYCMYCKCKAGMYNMCTCCAGCTCTYNTTCTCTCKKCCM 672

302 GACCACTTCAACATCAACCAATCAATGTTTCTCTCTCAATTAAGCTTGACTACTC 361
673 KACCKNTTCTGWSGCCCAKAYMYCAMMTCTCKAKSCCCYCCNNYCMNM 732

362 GACGCTGAAACATTAACCTTCAAGCTCTCATCTCTTATGCTTTCCAAACCGCTG 421
733 CWCMTCTMNAKCANCTTCTCTCTCMMTMAKCKWNNCTCCGSGACTYCTTAC 792

422 CTCTCATCCCAAGAAAGTTGATCATCTCTCACTGACATCAACGATTCAGTTC 481
793 TKMKCKNTCTCTTMCCTTMMCNCTCCMKNCCCTCCMNTCKTCTCTCNCMKRYC 852

482 GCAACAAACATGACCTGCTCTCTTGCACTGATGATGCACTGATCTCTTCAACA- 540
853 YVAKAKANCMCTCCCAKMKAKCTKCTCCCAKMSAKCKCCMCCCTCTATCWC 912

541 TCTTATATATCTCTTACTGACACGCGGCTGCTGCTCTCAACATTTAAAG 600
913 TCTCMTATCTCTCTCMYCMYMKCANCKYAYYCANCTMMNMCMCNCTCTCT 972

601 CTCACCTGATTTGCAAGAAAGAGTGAAGCTCAACGCGCACTGACGCGTCTT 660
973 NYCTCWCACGCTYCKCKCTMKCNMYMCRNCTYRCTCKKCCNCRNCKRMCKCTMCT 1032

QY 661 CCTCTACACATTCACACTGACGACGAGTGTACCATCTCCACACCCGCT 711
DB 1033 CTCGMMKMTCCGCMCCATCTMKSSTCTGCMGNCMTCCCTCNKCCYNNTKCY 1083

RESULT 5

US-09-072-967-328
Sequence 328, Application US/09072967
Patent No. 6592877
GENERAL INFORMATION:
APPLICANT: Reed, Steven G.
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Dillon, David C.
APPLICANT: Campos-Neto, Antonio
APPLICANT: Houghton, Raymond
APPLICANT: Vedvick, Thomas S.
APPLICANT: Iwardzik, Daniel R.
APPLICANT: Lodes, Michael J.
APPLICANT: Hendrickson, Ronald C.
TITLE OF INVENTION: COMPOUNDS AND METHODS FOR IMMUNOTHERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF TUBERCULOSIS
NUMBER OF SEQUENCES: 355
CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED AND BERRY LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
CITY: Seattle
STATE: Washington
COUNTRY: USA
ZIP: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/072,967
FILING DATE: 05-MAY-1998
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: MAKI, David J.
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 210121.411C9
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
TELEFAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 328:
SEQUENCE CHARACTERISTICS:
LENGTH: 1166 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: Genomic DNA
US-09-072-967-328

Query Match 2.0%; Score 63.8; DB 4; Length 1166;
Best Local Similarity 26.7%; Pred. No. 3.2e-08;
Matches 190; Conservative 136; Mismatches 384; Indels 1; Gaps 1;
QY 2 TCATATATTCGACCTCTCTTCATCTTCATCCAAAGACACAAAGCCGCATCG 61
DB 373 KSAAMTSMGSGSTCTMTYCNNGASATMTNMCCCGGAYVCSKNAVCCCTGTGCA 432
QY 62 CTTTCCCTCAACATCTCACTGACGACGCGCCGCTCTCTCACTTAAGGCTTCAATCG 121
DB 433 TYCCMCMGSGSGCTCCAMNCCACCTGNGYCCCTCCMCMTCYCATTCMNTCCGCTGCC 492
QY 122 TCTCATGCGCATCTCAACATATCGACCTGCGATATCATGAGCTGCGCTTTC 181
DB 493 TTTTMMCCSNCRTYCTCAMCNCSTKSGKACCNATMTCSACGHTCTMTMCSKAKMT 552
QY 182 ACCGCTCTTCATCTACCGCTGCTCCTCTCTCTCCAGAAAGAACTGAGCTCTCA 241
DB 553 TCCCTCNCCTYTNNCAMGMSCTCTMTMACTCKCCGGYCKNCKMYCTCTGCAV 612

QY 242 TTTCACTACAGACTTACACCAAGCCGGCTTACCACTTCTAGCTTTAAACCATC 301
DB 613 NMAACCKTCTYTCWMMYCMYCKCKAGWYKJMTCTCMACTGMYTTTCTCTCNKCCGA 672
QY 302 GACCACTTACACATCAACATCAATGATGTTTCTCTTCATTAAGCTTGACATAC 361
DB 673 KACCKNTTCTCWCSCCCCAKAKAYCYAMCMTMTCCMTCKACSCCCYCNMYCMMN 732
QY 362 GACCGCTGAACCTTATCACTCAAGCTCCCATCTCTCAATCGTTTCAACACCGCTG 421
DB 733 CWCMTCTCWNKAKCANCTTCTCTCTCMMYMTMAKCMGNNTCNCKSGACCTTCTAC 792
QY 422 CTCATCCCCCAAGAGCTTGTGCTCACTCTCACTGATCAACGAGTTCATGATCA 481
DB 793 TTKKCCKMTCTCTTCTMCKCTVMCMYCMKVNCCCTGMMTCTCTCTCNKNNRYCY 852
QY 482 GCAACCAACTGACCTGCTCTCTTGTGCACTCATATGACTCATCTCTCTCTCA 540
DB 853 YAKCAKCMCTCCCANMKAKCTCTCCCAKMKSAKNCCKCCCTCTATCCWC 912
QY 541 TCTTATCATCTCTCTTACTGACACCGTGGTCTCGCTCCACCATTCGATTTAAAG 600
DB 913 TCTCMTATCTCTCTCTCWCNMYCMYKMCACNCKCYAYTCACTMNMNCCANCTCTCT 972
QY 601 CTCACCTGATGTGCAAG 660
DB 973 NYTCMKRACGTYCKCKCKCTMCKNMYCMYKCTYRECTKCKCNCBANKMKMTCT 1032
QY 661 CCTCTACACATTCACACTGACGACGAGTGTACCATCTCCACACCCGCT 711
DB 1033 CTCGMMKMTCCGCMCCATCTMKSSTCTGCMGNCMTCCCTCNKCCYNNTKCY 1083

RESULT 6

US-08-232-463-14/c
Sequence 14, Application US/08232463
Patent No. 5670367
GENERAL INFORMATION:
APPLICANT: DORNER, F.
APPLICANT: SCHEIFLINGER, F.
APPLICANT: FALKNER, F. G.
TITLE OF INVENTION: RECOMBINANT FOWLPOX VIRUS
NUMBER OF SEQUENCES: 52
CORRESPONDENCE ADDRESS:
ADDRESSEE: Foley & Lardner
STREET: 1800 Diagonal Road, Suite 500
CITY: Alexandria
STATE: VA
COUNTRY: USA
ZIP: 22313-0299
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/232,463
FILING DATE:
CLASSIFICATION: 435
Prior Application Data:
APPLICATION NUMBER: US/07/935,313
FILING DATE:
APPLICATION NUMBER: EP 91 114 300.6
FILING DATE: 26-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 30472/114 IMMU
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 836-9300
TELEFAX: (703) 683-4109
TELEX: 899149

; INFORMATION FOR SEQ ID NO: 14:
 ; SEQUENCE CHARACTERISTICS:
 ; LENGTH: 7218 base pairs
 ; TYPE: nucleic acid
 ; STRANDEDNESS: single
 ; TOPOLOGY: linear
 ; IMMEDIATE SOURCE:
 ; CLONE: pTZapT-Fls
 ; US-08-232-463-14

Query Match 2.0%; Score 63.8; DB 1; Length 7218;
 Best Local Similarity 10.9%; Pred. No. 1.3e-07;
 Matches 53; Conservative 226; Mismatches 208; Indels 0; Gaps 0;

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QY 843 TATTACTACATATTAAAGTTTATCTTTAGTTTGGTCTGTTTATAGTTTCATCATGAG 902
DB 1551 TGTGACTAGCGTAGCATGCTCTTAGACGCATCTATTAGTTTCAAAAAAGCGCATGTG 1492
QY 903 TAAGTACGTGGAATAATCTCTATGAGACCCCTGATTTAAATGATGAAGCTAAGTCTTG 962
DB 1491 GCATCACTGTATTATCTATCTATGCAAGTAGTTAAAGATAGAAAGATTGTACRRR 1432
QY 963 GTCCACTAGACCGGAGTAGAGCAACATTTTACAGAGCTATAGGATGAATTGAGG 1022
DB 1431 RRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRR 1372
QY 1023 CTCTCAGCTCAGCTATCAAGAGAGCTGAAATCGCTAGAGGAAAGGCGCATGTC 1082
DB 1371 RRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRR 1312
QY 1083 GAGTGAATATGAGCTGATTGATGAGATATCAAAAGTATGAGCCAGAGTCAATGGC 1142
DB 1311 RRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRR 1252
QY 1143 CAAGAGACGAGCTACTAGCAAAATCCGAGAGCTTACAGTGAAGATATAGATT 1202
DB 1251 RRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRR 1192
QY 1203 CTTTGAGATGAATGACTTCTGGGAGACGATATCCCTGATATGACTTTAGCCCAT 1262
DB 1191 RRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRR 1132
QY 1263 GGGGTTACTGAGAGACGTCAGCATCTGTTGAGAAATGTCATCTGATAGAGAGACA 1322
DB 1131 RRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRR 1072
QY 1323 ATCGAGT 1329
DB 1071 RRRRRAT 1065

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RESULT 7
 US-09-949-016-16945
 ; Sequence 16945, Application US/09949016
 ; Patent No. 6812339
 ; GENERAL INFORMATION:
 ; APPLICANT: VENTER, J. Craig et al.
 ; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
 ; FILE REFERENCE: C1001307
 ; CURRENT APPLICATION NUMBER: US/09/949,016
 ; CURRENT FILING DATE: 2000-04-14
 ; PRIOR APPLICATION NUMBER: 60/241,755
 ; PRIOR FILING DATE: 2000-10-20
 ; PRIOR APPLICATION NUMBER: 60/237,768
 ; PRIOR FILING DATE: 2000-10-03
 ; PRIOR APPLICATION NUMBER: 60/231,498
 ; PRIOR FILING DATE: 2000-09-08
 ; NUMBER OF SEQ ID NOS: 207012
 ; SOFTWARE: FastSeq for Windows Version 4.0
 ; SEQ ID NO 16945
 ; LENGTH: 37155
 ; TYPE: DNA

; ORGANISM: Human
 ; US-09-949-016-16945

Query Match 1.9%; Score 59.8; DB 4; Length 37155;
 Best Local Similarity 47.7%; Pred. No. 7.1e-06;
 Matches 240; Conservative 0; Mismatches 257; Indels 6; Gaps 2;

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QY 89 CCGCCGCTCTCTCACTTACTGAGCTTCAATGCTCTCAATGCCATCTCTCAATCTG 148
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QY 149 ACCTGGCATATCACTGAGCTGCGGCTTCTCAACGCTCTCAATGCTACCGCTCT 208
DB 25894 CCTTACCCCTCAACCCCTCACTCAACCCCTCACTCACTCACTCACTCAACCCCTCACTCT 25953
QY 209 CCGCTCTCAAGGAAACATCGAGCTCTCAATTTCACTGACCTGACCTGACCAACAA 268
DB 25954 CACTACCCCTCACTCACTCTCAACCAACCCCTCACTCACTCACTCAACCCCTCACTCT 26013
QY 269 GCCGCTTCAACCACTTCTAGCTCTTAACTGACCTGACCACTTCAACCAATCAAA 328
DB 26014 CAC---TCACCCCTCACTCTCACTCACTCACTCACTCACTCACTCACTCACTCACT 26069
QY 329 TCGTTTCTCTCAATTAAGCTTGAATATCTGACCGCTGAACACTTATACCTTCAAG 388
DB 26070 TCTCTACTCATGCTCACTCACTCACTCACTCACTCACTCACTCACTCACTCACTCACT 26129
QY 389 CTCTCATCTCTTCACTGTTTCAACACCGCTGCTCTGATCCCCCAAGAGTTGTGAT 448
DB 26130 TACCCCTCACTGCTCAATACCCCTCACTGCTCAATACCCCTCACTGCTTCACT 26189
QY 449 CACCTGCTCATCACTCACTGATCTGATTCAGCAACCAACTGACCTGCTCTCTT 508
DB 26190 CAC--CCCTCACTCACTCACTCACTCACTCACTCACTCACTCACTCACTCACTCACT 26247
QY 509 GCACTATATGCTCATCTCTCTCAATCTTCAATCTTATCTTCTTATCTGACCAAC 568
DB 26248 TCACCCCACTCACTCACTCACTCACTCACTCACTCACTCACTCACTCACTCACTCA 26307
QY 569 GTGCTGCTCTCACTCACTGCTCC 591
DB 26308 CCCCTCACTCTCTCACTCACTCACT 26330

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RESULT 8
 US-08-658-136-2
 ; Sequence 2, Application US/08658136
 ; Patent No. 6071717
 ; GENERAL INFORMATION:
 ; APPLICANT: KLINGER, KATHERINE W
 ; APPLICANT: LANDES, GREGORY M
 ; APPLICANT: BURN, TIMOTHY C
 ; APPLICANT: CONNORS, TIMOTHY D
 ; APPLICANT: DACKOWSKI, WILLIAM
 ; APPLICANT: GERMINO, GREGORY
 ; APPLICANT: QIAN, FENG
 ; TITLE OF INVENTION: POLYCYSTIC KIDNEY DISEASE GENE
 ; NUMBER OF SEQUENCES: 58
 ; CORRESPONDENCE ADDRESS:
 ; ADDRESSEE: GENZYME CORPORATION
 ; STREET: ONE MOUNTAIN ROAD
 ; CITY: FRAMINGHAM
 ; STATE: MASSACHUSETTS
 ; COUNTRY: USA
 ; ZIP: 01701
 ; COMPUTER READABLE FORM:
 ; MEDIUM TYPE: Floppy disk
 ; COMPUTER: IBM PC compatible
 ; OPERATING SYSTEM: PC-DOS/MS-DOS
 ; SOFTWARE: PatentIn Release #1.0, Version #1.25
 ; CURRENT APPLICATION DATA:
 ; APPLICATION NUMBER: US/08/658,136
 ; FILING DATE:

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? CLASSIFICATION: 435
?
? ATTORNEY/AGENT INFORMATION:
?
? NAME: LASSEN, ELIZABETH
? REGISTRATION NUMBER: 31,845
? REFERENCE/DOCKET NUMBER: GEN4-17.8
? TELECOMMUNICATION INFORMATION:
?
? TELEPHONE: 508-872-8400
?
? TELEFAX: 508-872-5415
?
? INFORMATION FOR SEQ ID NO: 2:
?
? SEQUENCE CHARACTERISTICS:
?
? LENGTH: 53526 base pairs
?
? TYPE: nucleic acid
? STRANDEDNESS: single
? TOPOLOGY: linear
?
? MOLECULE TYPE: DNA (genomic)
?
? US-08-658-136-2

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Query Match	1.8%;	Score 55;	DB 3;	Length 53526;
Best Local Similarity	46.0%;	Pred. No. 0.0027;		
Matches 225; Conservative	0;	Mismatches 260;	Indels 4;	Gaps 1;

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Db	35885	CTCTCTCCCTCTCTTTTTCCTTCCCTTCCCTCCCTCCCTCTTCCCTCCCTCCCTCTTCCCTCCCT	35944
Qy	156	GATATTCATCGAGCTCGCGCTTTTACACGCTCTCTCATGTATGATACGGCTGCTCCCTCTC	215
Db	35945	TCCCTCTCTCTTCCCTCCCTCCCT	36004
Qy	216	TCCAAGGAACAATCGAGCTCTGCATTTTCATCTACCTACGACCTTCTACCAAGCGAGCT	275
Db	36005	TCCCTCTCCCTCTTCTCTCCCT	36066
Qy	276	TCACTACTTCTAGCTCTTAAACACTGACGACCTTACGATCAACCAATCAATGCTTTT	335
Db	36065	TTCCT	36124
Qy	336	CTCCCTCAATTAAGCTTGAACATCTGACGCGGTGAACATTTATCACTTCAAGCTCTCTGA	395
Db	36125	CT	36184
Qy	396	TCTCTTCACTGTTTTCACAACCGCTGCTCTCATATCCCTCCAGAAAGCTTGTGATCACTCT	455
Db	36185	TCT	36244
Qy	456	CACATCATCACAGTTCACTGATTCAGACCAACAACTCGACGCTGCTCTCTTGACATCTC	515
Db	36245	CTTCT	36304
Qy	516	ATATGTCATCTGATCTCTCTCTACCACTTTCATCATCTCCCTTACTGACACACGATGCTCTC	575
Db	36301	TTCT	36360
Qy	576	TTCGCTTCAC 584	
Db	36361	CCGCTTCCC 36369	

RESULT 9
US-08-658-136-1
; Sequence 1, Application US/08658136
; Patent No. 6071717
; GENERAL INFORMATION:
; APPLICANT: KLINGER, KATHERINE W
; APPLICANT: LANDES, GREGORY M
; APPLICANT: BURN, TIMOTHY C
; APPLICANT: CONNORS, TIMOTHY D
; APPLICANT: DACKOWSKI, WILLIAM
; APPLICANT: GERMINO, GREGORY
; APPLICANT: QIAN, FENG
; TITLE OF INVENTION: POLYCYSTIC KIDNEY DISEASE GENOME
; NUMBER OF SEQUENCES: 58
; CORRESPONDENCE ADDRESS:

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10 ADDRESS: GENZYME CORPORATION
11 STREET: ONE MOUNTAIN ROAD
12 CITY: FRAMINGHAM
13 STATE: MASSACHUSETTS
14 COUNTRY: USA
15 ZIP: 01701
16
17 COMPUTER READABLE FORM:
18 MEDIUM TYPE: Floppy disk
19 COMPUTER: IBM PC compatible
20 OPERATING SYSTEM: PC-DOS/MS-DOS
21 SOFTWARE: PatentIn Release #1.0, V
22
23 CURRENT APPLICATION DATA:
24 APPLICATION NUMBER: US/08/658,136
25
26 FILING DATE:
27
28 CLASSIFICATION: 435
29
30 ATTORNEY/AGENT INFORMATION:
31 NAME: LASSEN, ELIZABETH
32 REGISTRATION NUMBER: 31,845
33 REFERENCE/DOCKET NUMBER: GEN4-17,6
34
35 TELECOMMUNICATION INFORMATION:
36 TELEPHONE: 508-872-5410
37 TELEFAX: 508-872-5415
38
39 INFORMATION FOR SEQ ID NO: 1:
40
41 SEQUENCE CHARACTERISTICS:
42
43 LENGTH: 53577 base pairs
44 TYPE: nucleic acid
45 STRANDEDNESS: single
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47 TOPOLOGY: linear
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49 MOLECULE TYPE: DNA (genomic)
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Query Match	1.8%	Score 55	DB 3	Length 53577
Best Local Similarity	46.0%	Pred. NO	0.00027	
Matches 225	Conservative	0	Mismatches 260	Indels 4
				Gaps 1

QY	CTCTCTACCTTACTCGGGCTTCAATCGCTCTCATCGCTCATCGCATCTCTCAACATCACTGACCTGAC	155
Db	CTCTCTCTCCCTTCTTTTCCCTCTTCCCTCTCCCTCTCTCTTCCCTCTCCCTCTCTTCCCTCTTCCCTCT	359338
QY	156 GATATCACTGAGCTGCGCGCTTCTGACCGGCTCTCATCGCTCATCGTCAACCGCTGCTCCCTCTC	215
Db	35940 TCCCTCTCTTTCCTCCCTCCCTCTCTTTCCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT	359992
QY	216 TCGAAGGAAACAATCGAGCTCTCCATTTCACTCACTGACCTTACCAACAAGCCGCT	275
Db	36000 TCCCTCCCTCTTCTCTCCCTCCCTCTCTCCCTCTTCCCTCCCTCTCTTCCCTCTTCCCTCTTCCCTCT	360588
QY	276 TCACCACTTTAGCTCTTAAACAAGCTGACGACCTTACCATCAACAACAATGCTTT	335
Db	36060 TTCCCTCCCT	361128
QY	336 CTCTCTCATTAAGCTTGAATCACTGACGCGGAAACAATTACCTTCAAGCTCTCA	395
Db	36120 CT	361797
QY	396 TCTCTCATCTGTTTCAACAACGCGCTGCTCATCTCCCAACAAGCTTCTTCACTCT	455
Db	36180 TCT	362339
QY	456 CACTCATCAACAATCACTGATTCAGCAACAACAATGACCTGCTCTCTTGGACATC	515
Db	36240 CT	362989
QY	516 ATAGTCACTGAGTCTCTCTCAACAATTCATATCTCTCTTAATGAGACAACGAGCTC	575
Db	36296 TTCTCTCTTCT	363555
QY	576 TCGCTCAC 584	
Db	36356 CCGCTTCC 36364	

RESULT 10

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US-09-949-016-12147/c
; Sequence 12147, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
; APPLICANT: VENTER, J. Craig et al.
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
; WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
; FILE REFERENCE: CL001307
; CURRENT FILING DATE: 2000-04-14
; PRIOR FILING DATE: 2000-04-14
; PRIOR APPLICATION NUMBER: 60/241,755
; PRIOR FILING DATE: 2000-10-20
; PRIOR APPLICATION NUMBER: 60/237,768
; PRIOR FILING DATE: 2000-10-03
; PRIOR APPLICATION NUMBER: 60/231,498
; PRIOR FILING DATE: 2000-09-08
; NUMBER OF SEQ ID NOS: 207012
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 12147
; LENGTH: 767677
; TYPE: DNA
; ORGANISM: Human
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)...(767677)
; OTHER INFORMATION: n = A,T,C or G
US-09-949-016-12147

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Query Match      1.6%; Score 51.4; DB 4; Length 767677;
Best Local Similarity 49.3%; Pred. No. 0.025;
Matches 257; Conservative 0; Mismatches 246; Indels 18; Gaps 4;

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QY 62 CTTTCCCTCACAACCTCTCACTGACACCGCCGCTCTCTCACTTACTGCGCTTACGCG 121
DB 532452 CTTCTCACTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 532393
QY 122 TCTCATGCGCATCTCTCAACATCTGACCTGCGATATCACTCGAGCTGCGCTTCTC 181
DB 532392 TCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 532333
QY 182 ACCGCTCTCATCTGACACGCGCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 241
DB 532332 TCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 532280
QY 242 TTTCACTCACTGACCTCTACACGACGCGCTTACACCATCTTACCTTAAACACTC 301
DB 532279 TCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 532220
QY 302 GACCACTTCACTCACTCAACATCAATGTTTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 361
DB 532219 CTTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 532160
QY 362 GACGCTGACACTTATGACCTTCAAGCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 421
DB 532159 TCTTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 532107
QY 422 CTTCTATCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 481
DB 532106 CTTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 532047
QY 482 GCAACCAACTGACCTGCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 541
DB 532046 TTTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 531991
QY 542 CTTTCACTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 582
DB 531990 CTTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 531950

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RESULT 11
US-09-949-016-17361/c
; Sequence 17361, Application US/09949016
; Patent No. 6812339

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; GENERAL INFORMATION:
; APPLICANT: VENTER, J. Craig et al.
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
; WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
; FILE REFERENCE: CL001307
; CURRENT FILING DATE: 2000-04-14
; PRIOR FILING DATE: 2000-04-14
; PRIOR APPLICATION NUMBER: 60/241,755
; PRIOR FILING DATE: 2000-10-20
; PRIOR APPLICATION NUMBER: 60/237,768
; PRIOR FILING DATE: 2000-10-03
; PRIOR APPLICATION NUMBER: 60/231,498
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; NUMBER OF SEQ ID NOS: 207012
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; SEQ ID NO 17361
; LENGTH: 767677
; TYPE: DNA
; ORGANISM: Human
; FEATURE:
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; LOCATION: (1)...(767677)
; OTHER INFORMATION: n = A,T,C or G
US-09-949-016-17361

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Query Match      1.6%; Score 51.4; DB 4; Length 767677;
Best Local Similarity 49.3%; Pred. No. 0.025;
Matches 257; Conservative 0; Mismatches 246; Indels 18; Gaps 4;

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QY 62 CTTTCCCTCACAACCTCTCACTGACACCGCCGCTCTCTCACTTACTGCGCTTACGCG 121
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QY 122 TCTCATGCGCATCTCTCAACATCTGACCTGCGATATCACTCGAGCTGCGCTTCTC 181
DB 532392 TCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 532333
QY 182 ACCGCTCTCATCTGACACGCGCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 241
DB 532332 TCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 532280
QY 242 TTTCACTCACTGACCTCTACACGACGCGCTTACACCATCTTACCTTAAACACTC 301
DB 532279 TCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 532220
QY 302 GACCACTTCACTCACTCAACATCAATGTTTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 361
DB 532219 CTTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 532160
QY 362 GACGCTGACACTTATGACCTTCAAGCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 421
DB 532159 TCTTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 532107
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DB 532106 CTTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 532047
QY 482 GCAACCAACTGACCTGCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 541
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QY 542 CTTTCACTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 582
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RESULT 12
US-09-949-016-13343/c
; Sequence 13343, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
; APPLICANT: VENTER, J. Craig et al.
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

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RESULT 13
US-09-949-016-12818/C
; Sequence 12818, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
; APPLICANT: VENTER, J. Craig et al.
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
; TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
; FILE REFERENCE: CL001307
; CURRENT APPLICATION NUMBER: US/09/949, 016

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RESULT 14
US-09-949-016-16420
; Sequence 16420, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
; APPLICANT: VENTER, J. Craig et al.
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
; TITLE OF INVENTION: WITH HUMAN DISEASE. METHODS OF DETECTION AND USES THEREOF
; FILE REFERENCE: CL0010107
; CURRENT APPLICATION NUMBER: US/09/949,016
; CURRENT FILING DATE: 2000-04-14
; PRIOR APPLICATION NUMBER: 60/241,755
; PRIOR FILING DATE: 2000-10-20

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;; PRIOR APPLICATION NUMBER: 60/237,768
;; PRIOR FILING DATE: 2000-10-03
;; PRIOR APPLICATION NUMBER: 60/231,498
;; PRIOR FILING DATE: 2000-09-08
;; NUMBER OF SEQ ID NOS: 207012
;; SOFTWARE: FASTSEQ for Windows Version 4.0
;; SEQ ID NO: 16420
;; LENGTH: 234884
;; TYPE: DNA
;; ORGANISM: Human
;; FEATURE:
;; NAME/KEY: misc_feature
;; LOCATION: (1)...(234884)
;; OTHER INFORMATION: n = A,T,C or G
US-09-949-016-16420

Query Match 1.6%; Score 49.8; DB 4; Length 234884;
Best Local Similarity 45.5%; Pred. No. 0.031;
Matches 258; Conservative 0; Mismatches 302; Indels 7; Gaps 2;

QY 10 TCGACCTCTTCTTCATCTTGCATCCAAAGACACAAAGCGCCATCGCTTCCCT 69
DB 29923 TCCCTCTCCCTCTTCTTTCAGCTCCGCCACTTCTCTCTCCCTCCCTCCCT 29982
QY 70 CACAACTCTACTGACACCGCGCGCTCTCTACTTACTGAGCTTATGCTCTGACG 129
DB 29983 CCCCCCTCTCTCCCTCCCTCCCTCCCTCCCTCCCTCCCTCCCTCCCTCCCT 30042
QY 130 CCATCTCTCAATACTGACCTGCGATATGACTGAGCTGCGCTTCTACGCGCTC 189
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QY 190 TCCATCTGACCGCGCTGCTCTCTCTCCAAAGAAACAATGAGCTCTCATTTCACTC 249
DB 30103 TCTCTCCCT 30162
QY 250 ACTGACCTCTACACCAAGCGGCTTACCACTTACTGCTTTTACCACTGACCACT 309
DB 30163 CCTCTCTAC-----CCTTGTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 30217
QY 310 TCACATCAACCAATCAATCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 369
DB 30218 CTGCT 30277
QY 370 AACCTTATCACTTCAAGCTCTCTATCTCTTCTCTCTCTCTCTCTCTCTCTCT 429
DB 30278 CCTCATTTTCT 30335
QY 430 CCCCAAGAAAGTTGTCTATCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 489
DB 30336 CT 30395
QY 490 ACTGACCTGCT 549
DB 30396 GCTTCTCACT 30455
QY 550 TCTCTCTTACTGACCAAGCGCTCT 576
DB 30456 TCT 30482

RESULT 15
US-09-949-016-12345

;; Sequence 12345, Application US/09949016
;; Patent No. 6812339

;; GENERAL INFORMATION:

;; APPLICANT: VENTER, J. Craig et al.

;; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

;; FILE REFERENCE: CL001307

;; CURRENT APPLICATION NUMBER: US/09/949,016

;; PRIOR FILING DATE: 2000-04-14

;; PRIOR APPLICATION NUMBER: 60/241,755

;; PRIOR FILING DATE: 2000-10-20
;; PRIOR APPLICATION NUMBER: 60/237,768
;; PRIOR FILING DATE: 2000-10-03
;; PRIOR APPLICATION NUMBER: 60/231,498
;; PRIOR FILING DATE: 2000-09-08
;; NUMBER OF SEQ ID NOS: 207012
;; SOFTWARE: FASTSEQ for Windows Version 4.0
;; SEQ ID NO: 12345
;; LENGTH: 86414
;; TYPE: DNA
;; ORGANISM: Human
;; FEATURE:
;; NAME/KEY: misc_feature
;; LOCATION: (1)...(86414)
;; OTHER INFORMATION: n = A,T,C or G
US-09-949-016-12345

Query Match 1.6%; Score 48.6; DB 4; Length 86414;
Best Local Similarity 46.3%; Pred. No. 0.034;
Matches 230; Conservative 0; Mismatches 264; Indels 3; Gaps 2;

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QY 74 ACTCTACTGACACACCGCGCTCTCTCTACTTACTGCGCTTCAATGCGCAT 133
DB 67860 TGTTCATCT 67919
QY 134 CTCTCAATATCTGACCTGCGGATATATCTGAGCTGCGGCTTCTACCGCTCTCA 193
DB 67920 CACGCTATCT 67979
QY 194 TCGTACCGCTGCT 253
DB 67980 TCT 68039
QY 254 GACCTTACCAACCAAGCGGCTTACCACTTCTAGCTTTTAACTGACCTGACCTTAC 313
DB 68040 CT 68097
QY 314 CATCAACCAATCAATCT 373
DB 68098 CTTCT 68156
QY 374 CTATCACTTCAAGCT 433
DB 68157 CT 68216
QY 434 AGAAAGTTGTCTATCT 493
DB 68217 CCT 68276
QY 494 GACCTGCT 510
DB 68277 TGTCTGCGCT 68293

Search completed: July 22, 2005, 01:04:48
Jed time : 489 secs

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QY 1087 AGATATGAGCTGATTTGATGAGAGATATCAAAAGTATGAGCCAGAGTCAATGCGCAG 1146
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QY 1147 GAGAGAGAGCTATGAAACAAATCCGACGAGTTACAGTGGAGAGATATTCAGATTTCTT 1206
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QY 1207 GAGATGAATGACTTCTGAGGAAAGAGATTCCTGTATGAGACTTTAGCCGAGTTGGG 1266
Db 8995 AAGCTGATGAGTCTGTATGACAGAGATTCCTTGTCTGACCTCACTTGACAACTCGGA 9054
QY 1267 TTAAGTGAAGAGCTGACAGATTTGTTCCAGAAAGTGCATCTG----- 1308
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QY 1309 -----ATTAAGAGAGAGACAAATCGAGTTTCTTTTCCACTGCAATGTGAATNG 1355
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QY 1416 GTGATGAACAGCGGTACCAATTTATGATCAAGAAATTTGAAGAACTGTTGGTTCCCT 1475
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QY 1716 ACAAAG 1775
Db 9535 ACTAAG 9594
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QY 1836 GGAAGACATATGAGTGGGTGGGCTGTGAGCGCAATTTCTGAAGTTTGTGAGAGTTCCGCTC 1895
Db 9655 GCGCTCTGTGATAGTGGGCTGTGAGCAAGATTTCTGATGATGATGATGATGATGATGATGATGAT 9714
QY 1896 AAGAGATGAGGTATGAGCAAG 1955
Db 9715 ATTTCGTGAG 9774
QY 1956 TCTGAGTTTGAACATGTTGGCGACTTTTCAAGCGCTACAGGTTGAGAGATTTCAATGATTTAGA 2015

Db 9775 CTGAGTTTGAACATGTTTGAAGATTTTCCAGAGTTCAGTTTGAAGCTCTTACAGACAG 9834
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Db 9835 AGAGCTATACATCTTCTTCTTCTGAG 9894
QY 2076 GACTTCAAGCTCGGCTTGAAGATCTTTATTTTGAAGGAGAGAGAGAGAGAGAGAGAGAGAGAG 2135
Db 9895 GATTTTAAAGCTGAGATTTGAG 9954
QY 2136 AGTCAACCGAG 2195
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QY 2784 ACTTCACATTAATTAATATATCATCTGTGATTTGT--TCTTATTTTGTTCAGTGA 2840
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QY 2901 GTTTGGGAG 2958
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QY 2959 TCTAAGGATGAG 3018
Db 10774 TCTAAGGATGAG 10833
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QY	187	GAAGTTTGGGAATTCGCGCTCAAGAAAGTAAGGTTTAGACACGAGAAATGATGGACTTGGCA	1933
Db	9725	GATGACTTTGGGAATTCGCGCTCAATTCCTGCTGGACTCAGCGCGGACAAATGATATGCA	9784
QY	1934	TCACCTTGGCGCGAATGATGATCTCTGAGTTTGCACATGTTGGCGACTTTCACCGCTACAG	1993
Db	9785	GCACCTTAGCTGACTGGGAAATTCCTGGAGTTTGGCAATGTTGAAGATCTCCACAGGTTTGG	9844
QY	1994	GTTCCAGCAATTCATCGATTAGAAATCGCAACATTTCTTTTCCCTGCATTGAGCTATGAG	2053
Db	9845	GTTTGAGCACTCTCAACACAGAGAGCTTAACATCTTCTCCAGGCAATGAGGTGCACAG	9904
QY	2054	GATTCCTGAGGGGACGAAACATTTGACTTCAGCCGCGGCTTGAAGATCTTTATTTGGAGGG	2113
Db	9905	GATATATGAGGGAGATTAACATTAATGATTTTAAAGCCTGAGATTGGACGCTCTACTATGAGAA	9964
QY	2114	CAGTCCCGCCCACTGAGAGAGATTAGTCAACCGAAGAGCTTACATTAAGAAATGTTGATGA	2173
Db	9965	CGCTCCCGCCCATTTGAGATAGAGATGATCTTCTTGAAGAAAGCT-----GCCTCGAGATGG	10015
QY	2174	GACATATGATATAGATAGAGGCGGAGTTTGAACAGAGCATGTATCATTTGATGAGCATAT	2233
Db	10016	GATGATAGAAATAGGCGGACATGAAAGTTTCGACATAGACATGTATCATTTGCTGTAACATGT	10075
QY	2234	ACCTCCAGCGGAGGAAAAAGCAAGAGTTTGAGCGAAAGCTCACAGGAAACAACAGCAAGCTGCA	2293
Db	10076	ACCTCCAGCGGAGGAGAGCAAGAGCTTGAATGAAAGCTCATMAAAATACATTAATTTGCA	10135
QY	2294	GAACTGGTGCAGAAACAGGATTAAGTTAACTGGCAAGTGCTCGAGGGCTTACAAAGTTTCT	2353
Db	10136	GAACTGGTGCAGAAACAGGACAAAGCAAGCAAGCTGATCGCAAGTG-----TTTCAAGCTTCT	10186
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QY	2414	GGACATGCTTTCCAGGAGATATGACG-----GCCGAGCTTACAGAGCAGAA	2461
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QY	2462	GATTCTGCAATGTCCTCCTCGAGGGAATTCATCATTTCGAGCCCTCGAATCTTAGGAAGATAG	2521
Db	10307	AGTCCCAATGTCCTCCTCGAGGGAATTCATCATTTCTGGAACACAGAGGAAGAG	10366
QY	2522	GAGAAAGCACTCACTCGATCTAGCAGCAGAGACAGACGACTTCTGAGCTCTGATGTT	2581
Db	10367	GAAAGCTTACACTCACTCGATCTAGCAGC-----AATATCAGGCTCATTTATCTCAGAGAGATC	10423
QY	2582	ACGCGACCGCGGTGCTGCGCGCAATAGAGAAGAGAGTCTGAGTATCTCAAGCGGTGC	2641
Db	10424	ACTCGACCGGCGTCTGCGCGCAGCAGAGAAAGAGAGTCTGAGTATCTCAAGCGGTGC	10483
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QY	2702	CGGTTCTGTAATGCTTGGAGCAATACACAGGACCATTTGATCAACAATTCGTTCAAT	2761
Db	10544	AGGTTCTGTAATGCTTGGAGCAATACAGGACCATTTGATCAACAATTCGTTCAAT	10603
QY	2762	ATTGCACTGAGATAGCGGCTCACTTCACTTAATTAATATATCATCTTGTGATTTGTT	2821
Db	10604	CTTGCAGCTGAGATAGCGGCTCACTTCACTTAATTAATATATCATCTTGTGATTTGTT	10663
QY	2822	CTT-----TATTTTGTTTTCAATTTGATTTGATTTGTTCTCTGATATCTTCCAGTTTATTTCA	2878
Db	10664	CTTCAATTTTGTTTCTGATTTGATTTGATTTGTTCTCTGATATCTTCCAGTTTATTTCA	10723
QY	2879	CACAGTGAAGTGTGATTTAAAGTTTGGGGAAGGCTCAGAGAA--STATGTTGCATTTGA	2936
Db	10724	CACAGTGAAGTGTGATTTAAAGTTTGGGGAAGGCTCAGAGAGTGTGTGCTGATTTGG	10783
QY	2937	TATATTTTAAAGTGTGATTTCACTTAAGGCAATGAAAAACCAAAAAAATTTAAAAATTT	2996

Dd		10784	TATATTTTAGCTACATTCACTTAAGCATGAAAAACC-AAAAAAAAATTGAATAATT	10842
Oy	--	2997	CAGAAAATGATTCAC---AAAAAAAAAGTGTCCATGATAGTGCAATTACATTAGATCA	3053
Dd		10843	CAGAAATGATTCACAAAAAAGAGTGTTCACTGATGGTGCATCACAATTAGATCG	10902
Oy		3054	AGCTTAGAGTTCATTTAGATTTGTGTCATATGCATATGGGGATTAATGATGATAGCC	3113
Dd		10903	AGCTTAAGTGTTCATTTAGATTTGATGTTCATATGCATATGGGGATGATGATGATGCC	10962
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Dd		10963	TTGTAAACA 10971	
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		US-10-315-515-121		
		; Sequence 121, Application US/10315515		
		; Publication No. US2003016190A1		
		; GENERAL INFORMATION:		
		; APPLICANT: Wright, David A.		
		; APPLICANT: Voytas, Daniel F.		
		; TITLE OF INVENTION: NUCLEIC ACIDS RELATED TO PLANT		
		; TITLE OF INVENTION: RETROELEMENTS		
		; FILE REFERENCE: 08411-031001		
		; CURRENT APPLICATION NUMBER: US/10/315,515		
		; CURRENT FILING DATE: 2002-12-10		
		; PRIOR APPLICATION NUMBER: US 60/339,060		
		; PRIOR FILING DATE: 2001-12-10		
		; NUMBER OF SEQ ID NOS: 168		
		; SOFTWARE: FastSeq for Windows Version 4.0		
		; SEQ ID NO 121		
		; LENGTH: 13894		
		; TYPE: DNA		
		; ORGANISM: Arabidopsis thaliana		
		US-10-315-515-121		
	Query Match	32.9%; Score 1027.4; DB 16; Length 13894;		
	Best Local Similarity	68.6%; Pred. No. 1,2e-274;		
	Matches 1620; Conservative 0; Mismatches 626; Indels 117; Gaps 10;			
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Dd		8630 TAAATTTGTTTCATTTGCTATCTTTTGAGACTTAACCTATTTACATTTGAGCTTTGAGTTCTTA	8689	
Oy		880 CTGTTTTAGCTTTCATCATGATGACTACTACAGTGAAGAATCCGATATGAGCACCTGATAT	939	
Dd		8690 AAATCCCTAACAGGAATTCATGAGCAATTACAGTGCGAGTTCTTGTGTATCTGTGCTAC	8749	
Oy		940 AATGTGATGAACTAATGCTCTGTGCTCACTAGACCCGAGTGAAGCAACATGTTTAACGAG	999	
Dd		8750 AACATGATGAGCAGAAATGCTATCTTTCAAAGCCAGAGAGAAACAGAGAAATATCGAA	8809	
Oy		1000 AGCTAATAGGATGAATTTGAACGCTCTGACGCTGACGTAAATCAAAAAGAGCTGAATTC	1059	
Dd		8810 AGTTTCAG-----AAGGAAAAGCTGAGATA	8833	
Oy		1060 GCTAAGAGAAAGAGGCGCATGTCGATAGATATATGAGCTATGATGAGATATCAAAAAT	1119	
Dd		8834 GCCCAGAGAAAGAGACCGATGAGAGAGAGCTATGAACTTATAGACGMAATCTTGAGAGAC	8893	
Oy		1120 GAGTATAGCCCAAGTCAATGATGCGSCAAGAGACGAAGCTACTGAAACAAATCCGACGAGTT	1179	
Dd		8894 GAATGATGCTCTTAACAGACTGCGACAGCTACCAAACTTCTGACAAAGCCCAKATATTG	8953	
Oy		1180 ACAATGAGAGATATATCAGANTCTTTGAGATGAATGACTTCTTGCGGAAACGAGTATCCC	1239	
Dd		8954 CTTGCTGAGGAATATATTGAGGCTTTTCAAGCTGAATGAGTCTGTATGACGACGAGTATCCT	9013	
Oy		1240 TGATATGAGACTTTAAACCCAGTTGGGGTACTGAGGAGAGTGCAGATCTGTTCCAGAGAG	1299	
Dd		9014 TGCTGACCTCACTTGACCAACTCGGATGTGTGAAGATGTTCGACCTGTGACCAAAAT	9073	

QY 1300 TGTCACTG-----ATAAGAGAGCAATCGAG 1328
| | | | | | | | | | | | | | | | | | | | | |
Db TGTCACTGCACTTGTATGGCTTATCCGATATGACATATGAGATGAGCAATACAA 9133
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QY 1329 TTTCTTCCACATCGCAAGTGGAAATGATAGGAGCTCACAGACTTTGAGTGATACC 1388
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| | | | | | | | | | | | | | | | | | | | | |
QY 1389 ATGGGCTTAGGCTTGTACGTTCTTAGTGATGAACAGCGGTACAGATTAGATCAAG 1448
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Db GAAGATTTGGATTTTGTGATTTTCTGTATGTCTATGATGATGAGTTATCAATCAAG 9253
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QY 1449 AAATTTGAAGAAGCTGTTGGTTTCCCTAGTGAAGAAAGGAAACCAACCCAGTTTGACAG 1508
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Db CGATTTGAAGAAGTGTGATTTTCCAGTGGAAAGGATTTAAAGCCAAAGTATGAAGA 9313
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QY 1509 GAAGGCTTAAGATTTTGGGCTACTATTTGGAAACATTCAGCTTAATTCAGCGGG 1568
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Db GAAGGCTTAAGATTTTGGGCTACTATTTGGAAACATTCAGCTTAATTCAGCGGG 9373
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QY 1569 TCCAAGAGCAACCAATCCGAGCTCTGTGATTCGCTACTTTACAGCGCTCGGTTGCAAT 1628
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Db TCCAAGAGCAATCAGATACGAGCCCTGTCTATCAGTACTTCCAGCGTTCTGTAGCCAA 9433
| | | | | | | | | | | | | | | | | | | | | |
QY 1629 GTTTTAACTCCAGGAGCTTACAGGCAACCGTGTAAACAGACATGAAGATAGAT 1688
| | | | | | | | | | | | | | | | | | | | | |
Db GTACTTACTCCCGAGATTTACAGGAGCTGTCACTAATCTGATATGAGATGATCCCA 9493
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QY 1689 TCAGGCTTATAGGATTTCCGCGCTTACAAAGAAAGATGTCCTGAGAGAAATCTT 1748
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Db ATGGCCCTCAAGAACTCTCCGCAAACTTAAATGATGATGCTTCCAGGATTAAGTC 9553
| | | | | | | | | | | | | | | | | | | | | |
QY 1749 AACGATCAACCAACGATATGCTCTGTGATCATCTGTGTGGTATACATGAAGTGGCG 1808
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Db AATGACACACTCTCTATATCTTCTGTGATCATCTGTGTGATTAACAAACTGGGCG 9613
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QY 1809 CTGACAAACGCAAGAAAGATTAAGAGAGCACTATGCTGTGGTGGCTGTGTGACGCA 1868
| | | | | | | | | | | | | | | | | | | | | |
Db GTACAGCAATTAACGCAAGAGAGCAAGAGCGCTGTGTGATGAGTGGTGTGACACT 9673
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QY 1869 ATTTGAAAGTTTGTGAGTTCGCTCAAGAAAGTATGCTTTCAGCAACGAGATATGAC 1928
| | | | | | | | | | | | | | | | | | | | | |
Db ATTTGAAAGTTTGTGAGTTCGCTCAAGAAAGTATGCTTTCAGCAACGAGATATGAC 9733
| | | | | | | | | | | | | | | | | | | | | |
QY 1929 TTGATCACTTGGCGGAGTGTGATTTCTGTGATTTGACATGTTGGGAGCTTTACCGC 1988
| | | | | | | | | | | | | | | | | | | | | |
Db ATCGAGCACTTACCTGCTGCTGCAATTTCTGTGATTTGCAATGTTGGGAGCTTTACCGC 9793
| | | | | | | | | | | | | | | | | | | | | |
QY 1989 TACAGTTTCAGCATTTCAATTAAGAAATCCGCAATTTCTTTCCCTGCAATTAAGCT 2048
| | | | | | | | | | | | | | | | | | | | | |
Db TTCAGTTTTCAGCATTTCAATTAAGAAATCCGCAATTTCTTTCCCTGCAATTAAGCT 9853
| | | | | | | | | | | | | | | | | | | | | |
QY 2049 ACTAGATTTTCAGGAGCAAGCAATTTGACCTTCAAGCTCGCTTGAAGATTTTATTTTC 2108
| | | | | | | | | | | | | | | | | | | | | |
Db ACACGATTAATTCAGGAGCAAGCAATTTTGAAGCTTCAAGCTCGCTTCAATTTAT 9913
| | | | | | | | | | | | | | | | | | | | | |
QY 2109 GAGGCACTCCGCACTGAGAGATTAAGTCAACCGAAGAGACTACATATAGAAATGTT 2168
| | | | | | | | | | | | | | | | | | | | | |
Db GAGGCACTCCGCACTGAGAGATTAAGTCAACCGAAGAGACTACATATAGAAATGTT 9964
| | | | | | | | | | | | | | | | | | | | | |
QY 2169 GATGAGCATATGATATATGATAGGCGGAGTTTGAACAGAGCATATTCATTTCAATGAG 2228
| | | | | | | | | | | | | | | | | | | | | |
Db GATGAGCATATGATATATGATAGGCGGAGTTTGAACAGAGCATATTCATTTCAATGAG 10024
| | | | | | | | | | | | | | | | | | | | | |
QY 2229 CATATACCTCCAGCGCAAGAAAGCAAGTTTGAAGAGCTCACAGAAACAGCAAG 2288
| | | | | | | | | | | | | | | | | | | | | |
Db CATATACCTCCAGCGCAAGAAAGCAAGTTTGAAGAGCTCACAGAAACAGCAAG 10084
| | | | | | | | | | | | | | | | | | | | | |
QY 2289 CTGCAAGATGGTGAAGAAACAGAGATTAAGTTACTCGCAAGTGGCTGAGGCTATCAAG 2348
| | | | | | | | | | | | | | | | | | | | | |
Db TTGCAAGATGGTGAAGAAACAGAGATTAAGTTACTCGCAAGTGGCTGAGGCTATCAAG 10085
| | | | | | | | | | | | | | | | | | | | | |

QY 2349 TTTCTGAAGACAAAGATCAGCTGCTCCTTTCACATCACTATTTCCGCAATGACAGCTC 2408
| | | | | | | | | | | | | | | | | | | | | |
Db TTTCTGAAGACAAAGATCAGCTGCTCCTTTCACATCACTATTTCCGCAATGACAGCTC 10136
| | | | | | | | | | | | | | | | | | | | | |
QY 2409 CCTCAGGACATGCTTGGAGGATATGAGC-----GCCGAGCCTAGAGAG 2456
| | | | | | | | | | | | | | | | | | | | | |
Db CCTATGGAATGCCATGAGGAGAAATTAATCACTCGCGCACAGGCTTGAAGCTTATGAGCAG 10255
| | | | | | | | | | | | | | | | | | | | | |
QY 2457 CAGAAATTTTCATATGTCCTCGAGAGGATTCATATTCAGAGCTGTGAATTTAGAAAG 2516
| | | | | | | | | | | | | | | | | | | | | |
Db CAGAAATTTTCATATGTCCTCGAGAGGATTCATATTCAGAGCTGTGAATTTAGAAAG 10256
| | | | | | | | | | | | | | | | | | | | | |
QY 2517 AATGAGAGACGACATCTCATCTGATTTAGACAGAGACAGACGACATTTCTGAGTCTGT 2576
| | | | | | | | | | | | | | | | | | | | | |
Db AATGAGAGACGACATCTCATCTGATTTAGACAGAGACAGACGACATTTCTGAGTCTGT 10316
| | | | | | | | | | | | | | | | | | | | | |
QY 2577 AGTTTACGCGACCGCGGTGCTGGCGCAATGAAGAGAGGTGAGTATCTCAGAGC 2636
| | | | | | | | | | | | | | | | | | | | | |
Db AGTTTACGCGACCGCGGTGCTGGCGCAATGAAGAGAGGTGAGTATCTCAGAGC 10373
| | | | | | | | | | | | | | | | | | | | | |
QY 2637 GGTGCTGGCCGCAACAGAGCTGATGAGTACCAACCAATGCTGAGCTGATACGAA 2696
| | | | | | | | | | | | | | | | | | | | | |
Db GGTGCTGGCCGCAACAGAGCTGATGAGTACCAACCAATGCTGAGCTGATACGAA 10433
| | | | | | | | | | | | | | | | | | | | | |
QY 2697 CATGCGGTTGTCTATGCTTTGGAGCAATCAAGCAGCATTTGACTACCACTTGT 2756
| | | | | | | | | | | | | | | | | | | | | |
Db CATGCGGTTGTCTATGCTTTGGAGCAATCAAGCAGCATTTGACTACCACTTGT 10493
| | | | | | | | | | | | | | | | | | | | | |
QY 2757 TCATTTATTCGATGAGTGAAGGCTGACCTCACCATTAATTAATCATCTCTGTGAT 2816
| | | | | | | | | | | | | | | | | | | | | |
Db TCATTTATTCGATGAGTGAAGGCTGACCTCACCATTAATTAATCATCTCTGTGAT 10553
| | | | | | | | | | | | | | | | | | | | | |
QY 2817 TTGTTCTT---TATTTGTTTCAGTGATTTGATTTGCTGAGTACTCTTTCAAGTTT 2873
| | | | | | | | | | | | | | | | | | | | | |
Db TTGTTCTTCTTTTGTGTTTCTGTGATTTGATTTGCTGAGTACTCTTTCAAGTTT 10613
| | | | | | | | | | | | | | | | | | | | | |
QY 2874 ATTCAACAGTGAAGTGTGATTTAAGTTGGGAGAGGCTCAGAGAA--GTAATTTCA 2931
| | | | | | | | | | | | | | | | | | | | | |
Db ATTCAACAGTGAAGTGTGATTTAAGTTGGGAGAGGCTCAGAGAAAGTGTGTTGCA 10673
| | | | | | | | | | | | | | | | | | | | | |
QY 2932 TTGTATATATTTTAAAGCTGATTCATCTAAGGATTAAGAAACCAAAAAA----- 2986
| | | | | | | | | | | | | | | | | | | | | |
Db TTGTATATATTTTAAAGCTGATTCATCTAAGGATTAAGAAACCAAAAAA----- 10733
| | | | | | | | | | | | | | | | | | | | | |
QY 2987 -----TTAAATTTTCAAGAAATGATTTAC-----AAAAAAGTGTTCATGTTGAT 3039
| | | | | | | | | | | | | | | | | | | | | |
Db ATTTTGAATTTTCAAGAAATGATTTACAAAAAATGTTTCAATGTTGAT 10793
| | | | | | | | | | | | | | | | | | | | | |
QY 3040 TACATTTAGATCAAGTCTAAGTGTTCATTTAGATTTGATGATGATCAATAGGAGATA 3099
| | | | | | | | | | | | | | | | | | | | | |
Db CACATTTAGATCAAGTCTAAGTGTTCATTTAGATTTGATGATGATCAATAGGAGATA 10853
| | | | | | | | | | | | | | | | | | | | | |
QY 3100 ATGATGATAGCTTGTAAACA 3122
| | | | | | | | | | | | | | | | | | | | | |
Db ATGATGATAGCTTGTAAACA 10913
| | | | | | | | | | | | | | | | | | | | | |

RESULT 5
US-10-315-515-122
; Sequence 122, Application US/10315515
; Publication No. US20030166190A1
; GENERAL INFORMATION:
; APPLICANT: Wright, David A.
; APPLICANT: Voytas, Daniel F.
; TITLE OF INVENTION: NUCLEIC ACIDS RELATED TO PLANT
; TITLE OF INVENTION: RETROELEMENTS
; FILE REFERENCE: 08411-031001
; CURRENT APPLICATION NUMBER: US/10/315,515
; PRIOR APPLICATION NUMBER: 2002-12-10
; PRIOR FILING DATE: 2001-12-10
; NUMBER OF SEQ ID NOS: 168
; SOFTWARE: FastSeq for Windows Version 4.0

```

: SEQ ID NO 122
: LENGTH: 13966
: TYPE: DNA
: ORGANISM: Artificial Sequence
: FEATURES:
: OTHER INFORMATION: synthetically generated construct
US-10-315-515-122

```

Query Match	32.9%	Score 1036.8	DB 16	Length 13966
Best Local Similarity	68.5%	Pred. No. 1.8e-274		
Matches 1620	Conservative	0	Mismatches 627	Indels 117, Gaps 10

Qy	819	TTTGAATGTCGTTTTGGTTTTGGCTATGCAATACAAACAATTAAGCTTATCTTGAGTTTCG	878
Db	8666	TTCAATCTGTTTCATCTGCTATCTTTGAGACTTAACCTATTGACATTTGAGCTTTGAGTTCT	87255
Qy	879	TCTGTTTTTAAGTTTCATCATGAGTAATCAAGTGAATAATCTCTATAGACCCCTGATTA	938
Db	8726	AAATCCCTACAGAGAAATCATGAGCAATTAACGTGGCAGTTCTTCTGTGTATCTGACTA	87855
Qy	939	TAAATGTGATGAAGCTAAGTCTCTGTCCTCACTAGACCGGATGAGACAACTGTTTACGA	998
Db	8786	CAACATGATGAGACAGAAATGCTCATCTTCAAGGCACAGAGAGAGAACAGAGAAATACGA	88454
Qy	999	GAGCTATPAGGAGATGAATTTGAACGCTCGACGCTCGAGTATCAAAAGAGGCTGAAT	1058
Db	8846	AAAGTTTCG-----AAAGAAAGCTGAGAT	8865
Qy	1059	CGCTAAGAGAAAGAGGCGATGTCGATGAGATATAGCTGATGTGATGAGATTAACAAC	1118
Db	8870	AGCCCGAGGAAGAGAGCGATGAGAGAGGTAATGACTTTATAGCAAGATCTTGAGGA	89229
Qy	1119	TGAGTATGAGCGCAGAGTCATGCGCGACAGAGAACGAACTACTGAACAATCCGACGAGT	1178
Db	8930	CGAGTACATGCTGGAACAGACTCGGACAGCTACCAAACTTCTGCAACAAGCCGACATAT	8989
Qy	1179	TACAGTGAAGAGATATACAGATTTCTTGAATGAATATGATCTTGGGGAACGAGATATCC	1238
Db	8990	GCCGTGTAAGGAATATATGTAGAGCTTTTCAACCTGATATAGTCTGTACACAGAGTATCC	9049
Qy	1239	CTGATATGAGACTTTAGCCGAGTTGGGGGTATCTGGAGACGTCGACGACTGTTCGAACA	1298
Db	9050	TTGCTCGACCTCACCTTGACAACTCGAATTTGGAAGATTTTCAGCACTGTACCAAG	9109
Qy	1299	GTCATCATCTG-----ATAAGAGAGACATACGA	1327
Db	9110	TTGTGCATCTGGAACCTTGATGGCTTATCCGATATGTACATATGAAGATGAGACAATACA	9159
Qy	1328	GTTTCTTTCCACATGCAAGTGAAGAAATGTATGAGGGACTCACAGACTTTGAGCTGAGTAC	1387
Db	9170	ATTCTCTCCACACTACAAAGTATGAGCTTCAACCAAGTATGACCTTCGTATGATGGATTCG	9229
Qy	1388	CATGGGCTTAAAGCTTCTTGACGTTCTTATGTGATGAAACAGCGGTACCAATTTAGATCAA	1447
Db	9230	TGAAGATTTGAGATTTCTTGCAATTTCTGTGTATGTATGATAGTACAGGTTATCAATCAA	9289
Qy	1448	GAAATTTGAAACACTGTTGGTTTCCCTAGTGAAGAAAGGAAACCAACCCAGGTTTGAACAG	1507
Db	9290	GCGATTTGGAAGATTTGTTGATTTTCCAGTGTGAACGGGATCTAAGCCAAAGTATGAAG	9349
Qy	1508	GGAAGAGCTTAAAGATTTGTGGGCTACTATTATGGGAACAATCAACCGCTAAACTGACGCG	1567
Db	9350	AGAAAGATTGAAAGACTTGTGGATCACATCGGCGAGCTCTGTATACGGTTGAATGCTTCAG	9409
Qy	1568	GTCACAGAGCAACCAATCCCGAGTCTGTGATTCGCTATCTTACGGCTCGGTTGCCAA	1627
Db	9410	GTCAAAGGACATCATGATACGAGGCCCTGTCTACAGTATCTTCCAGCCTTCTGTAGCCAA	9468
Qy	1628	TGTTTTTTACTCCAGGAGTCTACAGGACCGTGTCTTAAACAAGACATGAAGATGATAGA	1687
Db	9470	CGTACTTACTCTCCGAGAGATTACAGGGAGCTGTCACTAATCTGTATATGAGAGATGATGC	9522
Qy	1688	TTCAAGCGTTATAGGAGATTTCTCCGCTTACAAAGAGAAAGATGTCTGAGAGAGATCT	1747

Db	9530	AATGGCCCTCAAGAAACTCTCCGCAAACTAATAATGGCAATGTCCCTCCAGGGTGAAGT	9589
Qy	1748	TAAAGACTCCACACAGTAATGCTCTGTGGATCCATCTGTGTGGTACATGAAGTGGC	1807
Db	9590	CAATGACACACTCTCTCTATATCTTCTTGATCCATCTGTGTGGATACAAAACCTGGGC	9649
Qy	1808	GCTGACAAACGGCAAGAAAGAGTAAAGAGAGACATATGCGTGGGTGGGCTTGACCGC	1867
Db	9650	GGTCACGCAATTAACCGCAAGAGACACAGAGCGCTCTGTGCATATAGGTGGGTGTGCACC	9709
Qy	1868	AATTCCTAAAGTTTGTGTGAGTTCCGCTCAAGGAATGAGGTTAGCACCGAATGATGA	1927
Db	9710	TATTCGTATGCTTGTGTGAGTCCCACTCATTTCTGTGSACTCTGAGCCACGACATGSA	9769
Qy	1928	CTTGATCACTTCGCGCGAATGTGAGTTCTGTGATTTGAATGTTGGGCACTTTCACCG	1987
Db	9770	TATCGAGCACCTACGTCACTGACCAATTCCTGGAGTTTGCAATGTTGACATTCACAG	9829
Qy	1988	CTACAGGTTGAGCAATTCATCGATTGAAATCGCCAACTTCCTTTCCCTCGCATTTACG	2047
Db	9830	GTTTCAGGTTTGACACTCTACACACAGGAGAGCTAACATCTTCTCCCTAGGCCCTGAGT	9889
Qy	2048	TACTAGGATTTCTGAGGGGACGAACATTAATGACTTCAAGCCTGGCGCTTAAGATCTTATTT	2107
Db	9890	CACACGATATAGAGGGAGATTAACATTTGATTTTGAAGCTTGAGATTGACGCTCTACTA	9949
Qy	2108	CGAGGCGAGTCCGCAACTGAGAGATTAGTCACACCGAAGAGCTTAAATGAAGATGT	2167
Db	9950	TGAGAACGCTCCACACTTAATGAGGACGATCTTCTTGAAGAGCT-----GCTTC	10000
Qy	2168	TGATGACACATATGATATAGATGAGCGGAGTTTGACACGAGCATGTATCATTTACGTGA	2227
Db	10001	GGATGGGATGAAAGAAATGAGACCAATGAAGTTGCACTACAGCATGTATCACCTTGTCTGA	10060
Qy	2228	GCATATTCCTCCAGCGAGAAAAGCAAGTTGAGCGAAAGCTCACAGGAACAACAGAA	2287
Db	10061	ACATGTACTCTCCAGCGAGGAGGACAAAGCTTGACTGAAGCTCATTAACATTTACATGA	10120
Qy	2288	GCTGCAGAATGTGTGCAAGAAACAGATTAATTAATCTCGGCAAGTGCCTCAGGGCTATCAA	2347
Db	10121	ATTGCAAGATGGTGTGCAAGAGAGACAGACAGCTGATTCGCAAGT-----TTTCAA	10171
Qy	2348	GTTTCTGAAGACAAAGATCACTGCTCTCTCTTCCACTACATATTTCCGCAATGACAGCT	2407
Db	10172	GCTTCTGACAGACAAAGTGAAGTTGCTTCTCTCCACCACTGCTATTCACAGGTCAACC	10231
Qy	2408	CCCTCAGGAATGCTTTCAGAGGATATGACG-----GCCGAGCCTTAGANA	2455
Db	10232	TCTATGGAATCCATCTCAGGAGAAATTATGCACTGTGCGCACAGGCTCTGAGCTTAGCA	10291
Qy	2456	GCAGAAATCTCTGCATGTCCCTCCAGAGCAATTCATCATTCGAGCCTCGTAATCTAGAA	2515
Db	10292	GCAGAGAGTCCACATGTCCAGGCTTAGGCAATTCATTCGATTCCTCCGGAAACAAGAG	10351
Qy	2516	GAATAGGAGAACGACATCACTCGATCTAGCAGACAGAGACAGACGATTCGTGCAGTTG	2575
Db	10352	AAGAAGGAAGGCTAACATCACTCGATCTAGCAGC---AGATCACGCTCATTCACCTGAG	10408
Qy	2576	TAGTTTACGGGACCCGCGTGTCTGTGGCCGCAATGAAAGAGAGGTGAGATATCTCAAG	2635
Db	10409	GAGATCACTGACCGTGTGTCTGTGGCCGACAGAGAAAGAGATGTGAGATTTCTCCAGG	10468
Qy	2636	CGGTCGAGGCGGCAACAGACTGATGAGATCGAGTACCAATGTCTGAGAGCTGATTCGGA	2695
Db	10469	CGGTGTGCGGCGCAACAGACTATATAGGTCAAGTACCAATGTCTGTGAGAGCTGATTCGGA	10528
Qy	2696	ACATGCGGTTGTGTATAGCTTTGGAGCAATCAACGAGCAGCTTATGACTCAACATTCG	2755
Db	10529	ACAAGGAGGTTTGTCTATAGCTCTGTGGAGCAATGCGAGGACGACATTTGAAGACCACTACG	10588
Qy	2756	TTTCAATTTGACGTGAGGTAAAGGCTCACATTCATATATATCATGCTTGTGA	2815

Db 10589 TTCATCTTCGACGAGTAAGCCCTCACTTCAACATTGTAATTATACCGTCTGTTGTA 10648
Qy 2816 TTTGTTCTT---TATTTGTTTCAGTGAATGGATTGTCTCGAGTACTCTTCCAAATT 2872
Db 10649 TTTGTTCTTCCTTTTGTGTTTCTGTGAATGGATTGTCTCGAGTACTCTTCCAAATT 10708
Qy 2873 TATTTACACAGTGAAGCTGTGTGATTTAAATTGGGGAGGGCTCAGGAA--GTATGTTGC 2930
Db 10709 TATTTACACAGTGAAGCTGTGTGATTTAAATTGGGGAGGGCTCAGGAAAGTGTGTTGC 10768
Qy 2921 ATGTATATATTTTAAAGCTGTCATTCATCTTAAGGATAGAAAAACCAAAAAA----- 2986
Db 10769 ATTGTGTAATTTTGAAGTCTGCATTCATCTTAAGGATAGAAAAACCAAAAAA----- 10828
Qy 2987 -----TTAAAAATTCAGAAAAATGATTTCAC---AAAAAAGAGTTCATGTAGTTGCA 3038
Db 10829 AATTTTGAATAATTCAGAAAAATGATTTCACAAAAAATGTTTCATGTAGTTGCA 10888
Qy 3039 TTACATTTTGAATCAAGTCTTAAGTGTTCATTTAGGATTTGTGCATATGCTTAGGGAT 3098
Db 10889 TCACATTTAGAGATCGATCTTAAGTGTTCATTTAGGATTTGTGCATATGCTTAGGGAT 10948
Qy 3099 AATGATGATPAGCCCTTGTAGCA 3122
Db 10949 GATGATGATPAGCCCTTGTAGCA 10972

RESULT 6

US-10-315-515-124
; Sequence 124, Application US/10315515
; Publication No. US20030166190A1
; GENERAL INFORMATION:
; APPLICANT: Wright, David A.
; APPLICANT: Voytas, Daniel F.
; TITLE OF INVENTION: NUCLEIC ACIDS RELATED TO PLANT
; FILE OF INVENTION: RETROELEMENTS
; FILE REFERENCE: 08411-031001
; CURRENT APPLICATION NUMBER: US/10/315,515
; CURRENT FILING DATE: 2002-12-10
; PRIOR APPLICATION NUMBER: US 60/339,060
; PRIOR FILING DATE: 2001-12-10
; NUMBER OF SEQ ID NOS: 168
; SOFTWARE: FASTSEQ for Windows Version 4.0
; SEQ ID NO 124
; LENGTH: 13320
; TYPE: DNA
; ORGANISM: Arabidopsis thaliana
US-10-315-515-124

Query Match 32.9%; Score 1025.6; DB 16; Length 13320;
Best Local Similarity 68.9%; Pred. No. 3.7e-274;
Matches 1620; Conservative 0; Mismatches 619; Indels 113; Gaps 11;

Qy 825 TTTGCTTTTGGTTTTCATTAATCAATTAACGTTTATCTTGAAGTTTCGCTGTT 884
Db 7962 TGGTTTCATGTCTATCTTTGAGACCTTAACCTATTTGACATTTGAGTTTGAATCT 8021
Qy 885 TTTAGTTTCATCATGATGATCACTACAGTGAATAATCCTCTATGACCCGTATTAATGT 944
Db 8022 CTACGAGAAATCAATGAGCAATTAACAGTGGCAGTTCTTGTGTGATCCGACTACAACT 8081
Qy 945 GGATGAAGCTAAGTCTGTGTCATGACCGAGGTGAGAGCAACATGTTTACGAGGCTA 1004
Db 8082 GGATGAGACAGAGTGTGTCATCTTCAAGCCAGAGAGAGAAAGAGATATGAAAGCTT 8141
Qy 1005 TAGGATGATTTGAACGCTCTGACAGCTGACGTATCAAGAAAGAGCTGAATGCTAG 1064
Db 8142 CAG-----TAGGAAAGCTGAGATAGCCCG 8165
Qy 1065 AGGAAAGAGGCGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1124
Db 8166 AGGAAAGAGGCGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 8225

Qy 1125 TGAGCCAGATCATTAGGCGCAAGAGAGAACTACTGAACAAATCCGACGAGTTACAGT 1184
Db 8226 CATCCCTGAAACAGACTGCGAGAGCTACCAACTTCTCACAAGCCAGCATTTAGCCCTAC 8285
Qy 1185 GAGAGATATATCATGATTTCTTTCAGATGATGATGATGATGATGATGATGATGATGATGAT 1244
Db 8286 TGAGGATATATTTAGGCTTTTCAAGCTGATGATGATGATGATGATGATGATGATGATGAT 8345
Qy 1245 TGAGATCTTTAGCCAGTGTGGGTTTACGAGAGAGTGCAGATGATGATGATGATGATGAT 1304
Db 8346 GACCTCACTTGCACAACTCGGATTTGGAAGATGTTCAAGACTGTATACCAAGTTGTGCA 8405
Qy 1305 TCTG-----ATPAGAGAGCAATCGATTCT 1333
Db 8406 TCTGACACTCTGATGCTTATCCGTTATGATGATGATGATGATGATGATGATGATGATGAT 8465
Qy 1334 TTCCACTGCAAGTGAATAATGATGAGGAGCTCAGACAGCTTTGAGCTGATATCATGAGG 1393
Db 8466 CTCACGCTGCAAGTGAAGCTCTACAAAGTATACCTCTGATGATGATGATGATGATGATGAT 8525
Qy 1394 GTTAGGCTTCTTGAAGTCTTATGATGAAACAGCGGTACCAATTTAGATCAAGAAATT 1453
Db 8526 ATGGGATTTCTGCGATTTCTGTGTATGCTGATGATGATGATGATGATGATGATGATGATGAT 8585
Qy 1454 GGAAGAACTGTTGTTGTTTCCCTAGTGAAGAGGAAACCAACCCAGTTTGAACAGAGAA 1513
Db 8586 GGAATGATTTATTTGGCTTCCCTAGTGAAGAGGAAACCAAGATCTTAAGCCAAAGTATGAAGA 8645
Qy 1514 GCTTAAGATTTTGGGCTACTATTTGGAAACATCTACCGCTAAATCCGAGCGTCCAA 1573
Db 8646 GTTGAAGATCTTGTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 8705
Qy 1574 GAGCAACCAATTCGAGATCCCTGTGATTCGTTACTTTCAAGGCTCGGTTGCCAATGTTTT 1633
Db 8706 GAGCATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 8765
Qy 1634 TTACTCCAGGAGTCTACAGGACCCGTGTCTTAACAGACATGAAGATGATGATGATGATGAT 1693
Db 8766 CTACTCC-GAAGATTTACAGGAGCTGATCACTGATGATGATGATGATGATGATGATGATGATGAT 8824
Qy 1694 GCTTATGAGGATTTCCGCGCTTACAAAGAGAAAGTCCCTGAGAGAGATCTTAAAG 1753
Db 8825 CTTCAAGAACTTTTCCGCAACTTAAGATGATGATGATGATGATGATGATGATGATGATGATGAT 8884
Qy 1754 CTCACACAGTAAATGCTCTGTGATTCATCTGTGTGATGATGATGATGATGATGATGATGATGAT 1813
Db 8885 CACAGCTCTCTATATCTTCTGTGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 8944
Qy 1814 AAACGCAAGAAAGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1873
Db 8945 CAATTAACCGAAAGAGACAGAGCGCTGTGTGATGATGATGATGATGATGATGATGATGATGAT 9004
Qy 1874 GAAAGTTTGTGAGTCCGCTCAAGAAAGTATGAGGTTGCAACGGAATGATGATGATGATGATGAT 1933
Db 9005 GATAGCTGTGTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 9064
Qy 1934 TCATTTGCGCGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1993
Db 9065 GCACTTACGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 9124
Qy 1994 GTTGAGATTCATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 2053
Db 9125 GTTTGAGCACTTATGAGAGAAAGTATGATGATGATGATGATGATGATGATGATGATGATGAT 9184
Qy 2054 GATTTCTGAGGCGAGAACTTTGATCTTCAACCGCTGCGCTTGAAGATCTTTATTTTCAGGG 2113
Db 9185 GATATATGAGGAGATTAATGATTTTGAAGCTGAGGTTGAGAGCTCTTACTATGAGAA 9244
Qy 2114 CAGTCCGCAACTGAGAGATTTAGTACACCGAGAGCTCAATATGAGATGATGATGATGATGAT 2173
Db 9245 CGCTCCGCACTTATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 9295
Qy 2174 GACATATGATATGATGAGGCGAGTTTGAACAGAGATGATATCTTCACTGAGCATAT 2233

Db 145 ATGAGAGAGAGTATGATGATTAAGACGAAGATCTGAGAGACGAGTACATGCTTGAACAG 204
 QY 1138 TGGCGAGAGAGAGAGAGTACTGATCAAAATCCGACAGGTTACAGTGGAGAGATATATC 1197
 Db 205 ACTCGAGAGAGTCAAACTTCTGACACAGCCGACATATGCTGCTGGAGATATGTT 264
 QY 1198 AGATCTTTGAGATGATGATCTTCTGGGAAACGAGTATCCCTGATATGAGACTTTAGCC 1257
 Db 265 AGGCTTTTCAAGCTGATGATGATCTTGTAGACAGAGGATCTTGTGACCTCAGCTTGA 324
 QY 1258 CAGTTGGGTTTACTGAGAGAGCTGACAGATCTGTTGAGAGATGTCATCTG----- 1308
 Db 325 CAAGCTGGATTTGAGAGATGTTGACAGCTGATCAAGATTTGATCAATCGACACTTTG 384
 QY 1309 -----ATPAGAGAGAGCAATGAGTTTCTTTCACACTGCA 1346
 Db 385 ATGGCTTATCCGATATGATGATGATGAGATGAGCAATCAATTCCTCTCAGACTACAA 444
 QY 1347 GTGGAATGATGAGAGAGTCAAGACTTTGAGCTGATACATGAGGTTAGGCTTTCTTG 1406
 Db 445 GTAGAGCTTCAACAGGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 504
 QY 1407 AGCTTCTTATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1466
 Db 505 CGATTTTCTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 564
 QY 1467 GGTTCCTTATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1526
 Db 565 GATTTTCCAGTGGAAACGAGATCTTAAAGCAAGATGATGATGATGATGATGATGATGATGAT 624
 QY 1527 TGGGCTACTATTTGAGAAACAATCTACCGCTTAACTGACGCGGTGCAAGAGCAACAATC 1586
 Db 625 TGATATCAATCGGAGAGCTGATGATGATGATGATGATGATGATGATGATGATGATGAT 684
 QY 1587 CGGAGTCTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1646
 Db 685 CGGAGCTCTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 744
 QY 1647 TCTACAGGAGCTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1706
 Db 745 ATTACAGGAGCTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 804
 QY 1707 CTCGCGCTTCAAAAGAGAAAGATGATGATGATGATGATGATGATGATGATGATGATGAT 1766
 Db 805 CTCGCGCAAACTTAAATAGGATGATGATGATGATGATGATGATGATGATGATGATGAT 864
 QY 1767 ATGCTCTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1826
 Db 865 ATACTCTTCTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 924
 QY 1827 AAGGTAAAGAGAGAGCTATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1886
 Db 925 AGAGCAGAGGCGCTGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 984
 QY 1887 GTTCCGCTCAAGAGAGAGGTTAGACACGAGATGATGATGATGATGATGATGATGATGAT 1946
 Db 985 GTTCCACTCAATTTCTGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1044
 QY 1947 TGTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 2006
 Db 1045 TGGCAATTTCTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1104
 QY 2007 TCGATTAAGATGCGCAATTTCTTCTGATGATGATGATGATGATGATGATGATGATGAT 2066
 Db 1105 ACAGAGAGAGAGAGTCAATCTTCTGATGATGATGATGATGATGATGATGATGATGAT 1164
 QY 2067 AGGAACTTGAATCAAGCTGATGATGATGATGATGATGATGATGATGATGATGATGAT 2126
 Db 1165 GATTAACATTTATTTAGGCTGATGATGATGATGATGATGATGATGATGATGATGATGAT 1224
 QY 2127 GAGAGATTAAGTCAACAGAGAGAGTCAATTAAGAGATGATGATGATGATGATGATGAT 2186
 Db 1225 GATGAGAGAGATCTTCTTGAAGAGCT-----GCTTCGATGATGATGATGATGATGAT 1275

QY 2187 GATGAGCGAGAGTGTGACAGAGATGATGATGATGATGATGATGATGATGATGATGATGAT 2246
 Db 1276 GAGAGAGTAAAGTTGACACTGATGATGATGATGATGATGATGATGATGATGATGATGAT 1335
 QY 2247 AAAAGAGAGTGTGACAGAGTCAACAGAGACAGAGAGTCAAGAGTGTGACAG 2306
 Db 1336 CAGAGAGAGGCTTGTGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1395
 QY 2307 AAGAGAGTAAAGTAACTGAGAGTGTGATGATGATGATGATGATGATGATGATGATGATGAT 2366
 Db 1396 AAGAGAGAGAGTGTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1446
 QY 2367 AGTGTCTCTTCCACTCAATGATGATGATGATGATGATGATGATGATGATGATGATGAT 2426
 Db 1447 AGTGTCTCTTCCACTCAATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1506
 QY 2427 AGGAGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 2474
 Db 1507 AGGAGATTAATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1566
 QY 2475 CTTGAGAGAGTCAATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 2534
 Db 1567 CAGGCTAGAGCTTGTGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1626
 QY 2535 ACTGATCTTACAG 2594
 Db 1627 ACTGATCTTACAG 1683
 QY 2595 GCTGCGCGAGTAAAG 2654
 Db 1684 GCTGCGCGAG 1743
 QY 2655 GCTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 2714
 Db 1744 GCTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1803
 QY 2715 GCTTGGAGAGAGTCAAG 2768
 Db 1804 GCTTGGAGAGAGTCAAG 1857

RESULT 9

US-10-615-005-24
 ; Sequence 24, Application US/10615005
 ; Publication No. US20040016018A1
 ; GENERAL INFORMATION:
 ; APPLICANT: Wright, David A.
 ; APPLICANT: Voytas, Daniel F.
 ; TITLE OF INVENTION: Plant Retroelements and Methods Related Thereto
 ; FILE REFERENCE: F-1065 ISURF Plant Retroelement
 ; CURRENT FILING DATE: 2003-07-08
 ; PRIOR APPLICATION NUMBER: US/09/322,478
 ; PRIOR FILING DATE: 1999-05-28
 ; PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/087125
 ; PRIOR FILING DATE: EARLIER FILING DATE: 1998-05-29
 ; NUMBER OF SEQ ID NOS: 41
 ; SOFTWARE: PatentIn Ver. 2.0
 ; SEQ ID NO 24
 ; LENGTH: 1857
 ; TYPE: DNA
 ; ORGANISM: Arabidopsis thaliana
 ; US-10-615-005-24

Query Match 24.1%; Score 751.6; DB 17; Length 1857;
 Best Local Similarity 65.3%; Pred. No. 3.3e-198;
 Matches 1250; Conservative 0; Mismatches 564; Indels 100; Gaps 6;

QY 898 ATGATTAATCAAGTGAAGAAATCTTATGAGACCTGATTAATATGATGATGATGATGAT 957
 Db 1 ATGAGCAATTAACAGTGAAGTCTTCTGTTGATCTGATCAACAATGATGATGATGATGAT 60

QY 958 TCCTGCTCACTAGACCGGAGTGAGACAACTGTTTACGAGAGCTTATAGGATCAATTT 1017
Db 61 TCGTCATCTTCAAGGCCAGAGAGAAACAGAGAAATCGAAAGTTTCAG----- 110
QY 1018 GAACGCTCTGACGCTCGACGTAATCAAGAGAGAGCTGAATTCGCTAGAGAGAGGGCG 1077
Db 111 -----AMGAAAGCTGAGATAGCCCGAGAGAAAGAGAGG 144
QY 1078 ATGTCGAGTAGATATAGAGCTGATGATGAGATATCAAAACTGATATAGCCAGAGTCA 1137
Db 145 ATGAGAGAGAGGTATAGAGCTTATAGACAGAGATCTGAGAGAGAGATCACTGCTGAAACG 204
QY 1138 TGGCGCAAGAGAGACGAAAGCTACTGAACTAAATCCAGACAGGTTTACAGTGAAGAGATATC 1197
Db 205 ACTCCAGAGGTACCAAACTTCTGCACTAGCCGACATATTCCTGCTGAGGAAATATGTT 264
QY 1198 AGATCTTTGAGATGAATGACTTTCGGGGAAGAGGTATCCGATATGAGACTTTAGCC 1257
Db 265 AGGCTTTTCAAGCTGATATGATGTTCTGTAGACAGAGGTATCTTGCTCGACTTCAGCTTGCA 324
QY 1258 CAGTTGGGGTTTACTGAGAGACGTGACGATCTGTTGAGAAAGTGTATCTG----- 1308
Db 335 CAACCTCGATGTTGTGGAAGATGTTGACGACCTGTACCAAAAGTGTATCTGAGCACTTTG 384
QY 1309 -----ATPAGAGAGACAAATCGAGTTTCTTTCCACTGCA 1346
Db 385 ATGCTTATCCGTATGATAGCATATGAAATGAGACAAATTCCTCTCCACACTACAA 444
QY 1347 GTGGAATGATAGAGGACTCAACAGCTTTGAGCTGAGATACATAGGGGGTTAGGCTTTCTG 1406
Db 445 GTAGAGCTCTACCAAGATGAGACTCTGTAGTAGGTGATTTGAGAGATGGGATTCCTG 504
QY 1407 AGCTCTTATAGTAGAATGACAGCGGTACAGATTTAGATCAAGAAATTGGAAAGACTGTTT 1466
Db 505 CGATTTTCTGTATATGTCATATGATGATGATGATGATGATGATGATGATGATGATGATGAT 564
QY 1467 GGTTCCTCTAGTGAAGAGAAACCAACCCAGGTTTGAAGAGAGAGAGAGAGAGAGAGAGAG 1526
Db 565 GATTTTCCAGTGAACCGGATCTAAGCCAAAGTATGAAGAGAGAGAGAGAGAGAGAGAGAG 624
QY 1527 TGGGCTACTATTTGGGAAACAACTTACCCGCTAACTGACCGCGTCCAGAGGACCAACAAATC 1586
Db 625 TGGATCACCATGCGGCTCTGTACCGTTGAATGCTTCCAGGTCAAAGAGCAATCAGATA 684
QY 1587 CGAGTCTCTGATTCGCTACTTTCAGCGCTCGGTTGCCAATGTTTTCCTCAAGAGAG 1646
Db 685 CGGAGCCCTGTATCAGGTACTTCCAGGCTTCTGTAGCCAAAGTACTTACTCTCCGAGAG 744
QY 1647 TCTAAGGACACCGTGTCTAACAACAAGATGAAGATGATGATGATGATGATGATGATGAT 1706
Db 745 ATTACAGGGACTGTCACTAATCTGTATGTAGAGATGATGATGATGATGATGATGATGATGAT 804
QY 1707 CTCCGCTTACAAAAGAGAAATGCTCTGAGAGAGAGATCTTAAAGACTCACCAAGATA 1766
Db 805 CTCCGCGCAAACTAAAGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 864
QY 1767 ATGCTCTGTTGATCCATCTGTGTGGGTACATGAATGGGCGCTGCAAAAGGCAAGAG 1826
Db 865 ATACTTCTTCTGATCATCTGTGTGATACAAAAATGGGCGGTGCAATTAACCGCAAG 924
QY 1827 AAGTAGAGAGAGACTATGCGTGGGTGCGTGTGAGAGCCCAATTCYAAAGTTTGTGA 1886
Db 925 AAGAGACGAGGCGCTGTGCAATAGGTGGGTGTGACACTATTCYAAAGTTTGTGA 984
QY 1887 GTTCCGCTTAAAGAGTAGGGTTTGAACCGAAGATATGATGATGATGATGATGATGATGATGAT 1946
Db 985 GTTCCCACTAATTTCTGCTGAGACTCGAGCGACGCAATGATATGATGATGATGATGATGATGAT 1044
QY 1947 TGTGATTTCTGAGTTTGAATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 2006
Db 1045 TGTGCAATTTCTGAGATTTGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1104
QY 2007 TGTGATTAAGATGCAACATTTCTTTCCCTGTGATTAAGCTAATGATTTCTGAGAGGCT 2066

Db 1105 ACAGACAGAGAGGTATCAATCTTCTCTTACGCTGAGGTCACACGATATATGAGGGA 1164
QY 2067 AGGAATTTGACTTCAAGCCCTGCGCTTGAAGATCTTTATTTGAGGGGACGCCCAACT 2126
Db 1165 GATATCAATTTATTTAGGCTTGAAGATTTGAGAGGCTCTTACTATGAGAAAGCTTCAACATTA 1224
QY 2127 GAGGAGATTTAGTCAACCGAAGAGCTACATATGAGATGTTGATGAGATATGATATTA 2186
Db 1225 GATGAGAGCATTTCTTTGAAGAACT-----GCTTGGATGGGATGATGAAGAT 1275
QY 2187 GATGAGGCGGATTTGACACGAGATATGATATTTGAGTGAATATATCTTCCAGGAG 2246
Db 1276 GAGAGATTAAGTTTGAACACTAGATGATGATGATGATGATGATGATGATGATGATGATGAT 1335
QY 2247 AAAAGAGAGTTTGAAGAGAGCTCAAGAGAAACAGACAGAGCTGAGAGAGTGTGCAAG 2306
Db 1336 CAGACAAAGAGCTTGTACGAGAGCTCAATAGAAATTAAGTAAATTTGAGAAAGTGTGCAAG 1395
QY 2307 AAACAGATTAAGTTACTGCGCAAGTGTCTGAGGCTATCAAGTTTCTGAGAGACAGATC 2366
Db 1396 AAGCAGACAGAGTATGATGCAAGTG-----TTTCAAGCTTGTGACAGACAGAGCTG 1446
QY 2367 AGCTGCTCTCTTCCACTCACTAATTTCCGCAATGACAGCTCCCTCCAGAGATGCTTGG 2426
Db 1447 AGTGTCTTCTTCCACACGCTATTCACAGGTACAACTCTTATGAAATGCAATG 1506
QY 2427 AGGAGATATGACG-----GCCGAGCTTGAAGAGAGAGATTTCTGATGTC 2474
Db 1507 AGGAGATTAATGACACCTGCGCACAGGCTGTAGCTTACGAGAGAGAGATGCCAATGTC 1566
QY 2475 CTTGAGAGCATTCATATTCCTGAGCTCTGATATCTGAGAAATGAGAGAAACGACATC 2534
Db 1567 CAGGCTAGGACTTGTCTATTTGAAATCCCGGAGAACAAAGAGAAAGGATCACTC 1626
QY 2535 ACTGATTTAGCAGAGAGAGAGAGACTTTCGAGTGTGATGATTTACCGGACCGCGT 2594
Db 1627 ACTGATCTTACAGG---AGATCAAGCTCTCATTTCACTGAGAGAGATCACTGACCGTGT 1683
QY 2595 GCTGCGCGCAATTAAGAGAGAGAGAGTATCTCTGAGAGCGGTGCTGCGCACAGA 2654
Db 1684 GCTGCGCGCAGAGAGAGAGAGATGTCAGATTTCTCTGAGAGCGGTGCTGCGCACAGA 1743
QY 2655 GCTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 2714
Db 1744 GCTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1803
QY 2715 GCTTGGAGCAATCAG 2768
Db 1804 GCTTGGAGCAATCAG 1857

RESULT 10
US-10-395-607-24
; Sequence 24, Application US/10395607
; Publication No. US20040019928A1
; GENERAL INFORMATION:
; APPLICANT: Wright, David A.
; APPLICANT: Voytas, Daniel F.
; TITLE OF INVENTION: Plant Retroelements and Methods Related
; FILE REFERENCE: 08411/036001
; CURRENT APPLICATION NUMBER: US/10/395,607
; CURRENT FILING DATE: 2003-03-24
; PRIOR APPLICATION NUMBER: 09/586,106
; PRIOR FILING DATE: 2000-06-02
; PRIOR APPLICATION NUMBER: 09/322,478
; PRIOR FILING DATE: 1999-05-28
; PRIOR APPLICATION NUMBER: 60/087,125
; PRIOR FILING DATE: 1998-05-29
; NUMBER OF SEQ ID NOS: 200
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 24

LENGTH: 1857
 TYPE: DNA
 ORGANISM: Arabidopsis thaliana
 US-10-395-607-24

Query Match 24.1%; Score 751.6; DB 17; Length 1857;
 Best Local Similarity 65.3%; Pred. No. 3.3e-198;
 Matches 1250; Conservative 0; Mismatches 564; Indels 100; Gaps 6;

898 ATGAGTACTACAGTGGAAAAATCCTTATGACCCCTGATTATATGATGATGAAGTAA 957
 1 ATGAGCAATTACAGTGGAGTCTTCTGTGATCCGACTACCAATGATGAGAGAGA 60
 958 TCTCGTCCATAGACCGAGATGAGACAATGTTTACAGAGCTATAGGATGATTT 1017
 61 TGTGATCTTCAAGGCCAGAGAGAGACAAGAGATACGAAGTTTCAG----- 110
 1018 GAACGCTGACAGCTGACGTATCAAGAGAGCTGAATCGCTAGAGAGAGAGGCG 1077
 111 -----AGGAAAGCTGAGATAGCCGAGAGAAAGAGAGCG 144
 1078 ATGTGAGATGATATGAGCTGATTTGATGAGATATCAAACTGATATGAGCCAGTCA 1137
 145 ATGAGAGAGAGATGAGCTTATATGACGAAGATCTGAGAGACGAGTACATGCTTGAACG 204
 1138 TGGCGCAAGAGAGCAAGGCTACTGACCAATCCGACGAGGTTACAGTGGAGATATTC 1197
 205 ACTCGCAGAGCTACCAACTTCTGCAAGCCGACATTTGCTGCTGAGAAATGTT 264
 1198 AGATCTTGTAGATGATGATCTTCTGAGGAGACGAGATATCCCTGATATGAGCTTTAGCC 1257
 265 AGGCTTTTCAAGTGAATGATTTCTGTACAGAGGATATCCTTGTGACCTCATTGCA 324
 1258 CAGTTGGGTTACTGAGAGAGCTGACGATCTGTTGAGAGATGTTGATCTG----- 1308
 325 CAACCTGAGTGTGAGAGATGTTGAGACCTGTACCAAGTTGATCTGAGACCTTTG 384
 1309 -----ATAGGAGAGCAATGAGTGTCTTTCCACACTGCA 1346
 385 ATGGCTTATCCGATATGATATGAGATGAGAGCAATACATCTCTCCACACTCA 444
 1347 GTGGAATGTATGAGGAGACTCAGACTTTGAGCTGATACCATGAGGTTAGGCTTTG 1406
 445 GTAGAGCTTACCAAGTATGACCTGTGATGATTTGATGATGATGATGATGATGATG 504
 1407 AGCTTCTTAGTGAGTGAACGCGGATCCAGATTTAGATCAAGAAATTTGAGAGATGTT 1466
 505 CGATTTTCTGTGATGTGATGATGATGATGATGATGATGATGATGATGATGATGATG 564
 1467 GGTTCCTTCTAGTGAAG 1526
 565 GATTTTCCAGTGAAG 624
 1527 TGGGCTACTATTGGGAGCAATCTACCGCTAACTCGACGCGTCCAGAGCAACCAATTC 1586
 625 TGGATCACATCGGAGCTCTGACCGTTGATGCTTCCAGGCTCAAGAGCAATCGATG 684
 1587 CGGAGTCTGTGATGCTACTTTCAGCGCTCGGTTCCAAATGTTTTTACTCCAGGAG 1646
 685 CCAGGCTGTGATGAGTACTTCCAGCGTTCTGTAGCCACGTACTACTCCGAGAG 744
 1647 TCTACAGGAGCGGTGCTTACACAGACATGAGAGATATGATGATGATGATGATGATG 1706
 745 ATTACAGGAGCTGTCTACTCTCTGATATGAGAGATATGCAATGCGCCCTCAAGAGAACT 804
 1707 CTCGCGCTTACAAAGAGAAAGATGCTGAGAGAGATCTTAACTACCACTACCAAGTAA 1766
 805 CTCGCGCAAACTTAAATGAGATGCTCCAGGAGTGAATGATGATGATGATGATGATG 864
 1767 ATGCTCTGTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATG 1826
 865 ATACTCTTCTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATG 924

QY 1827 AAGTAAAGAGAGACTATGCTGGTGGCGCTTGTGACCCCAATCTTGAAGTTGTGGA 1886
 DB 925 AAGAGCAGAGCGCTCTGTGATATGAGTGGCGTGTGACACCTATCTGATAGCTTGTGA 984
 QY 1887 GTTCCGCTCAAGAGATAGGTTAGCACCGAAGATATGATGATGATGATGATGATGATG 1946
 DB 985 GTCCCACTATTTCTGTGATGATGATGATGATGATGATGATGATGATGATGATGATG 1044
 QY 1947 TGTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATG 2006
 DB 1045 TGCATTTCTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATG 1104
 QY 2007 TCGATTAAGTCCCAATCTTCTTCCCTGATTAAGCTATGATGATGATGATGATGATG 2066
 DB 1105 ACAGACAGAGAGCTTAACTATCTCTCCCTAGCCCTGAGGTCAACAGATATGAGGGA 1164
 QY 2067 AGGACATTTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATG 2126
 DB 1165 GATTAATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATG 1224
 QY 2127 GAGGAGATTAGTCAACCGAAGGCTACATTAAGATGATGATGATGATGATGATGATG 2186
 DB 1225 GATGAGAGAGATCTTCTTGAAGAGCT-----GCTTCGATGAGATGATGATGATG 1275
 QY 2187 GATGAGCGGAGTGTGACACGAGATGATATCAATTCAGTATGATGATGATGATGATG 2246
 DB 1276 GAGAGCATTAAGTGTGACATGAGATGATGATGATGATGATGATGATGATGATGATG 1335
 QY 2247 AAAAGCAGAGTTTGAAGCAAGCTCACAGAACACAGAGAGTGCAGAGTGTGCAAG 2306
 DB 1336 CAGAGCAGAGCTTGTGATGATGATGATGATGATGATGATGATGATGATGATGATG 1395
 QY 2307 AAAAGCAGATATGATGATGATGATGATGATGATGATGATGATGATGATGATGATG 2366
 DB 1396 AAGCAGAGAGAGCTGATGATGATGATGATGATGATGATGATGATGATGATGATG 1446
 QY 2367 AGTGTCTCTTCTTCACTCACTCACTCACTCACTCACTCACTCACTCACTCACTCACT 2426
 DB 1447 AGTGTCTCTTCTTCACTCACTCACTCACTCACTCACTCACTCACTCACTCACTCACT 1506
 QY 2427 AGGAGATATGAGC-----GCCGAGCTTGAAGAGAGAGAGAGAGAGAGAGAG 2474
 DB 1507 AGGAGATATGAGC-----GCCGAGCTTGAAGAGAGAGAGAGAGAGAGAGAGAG 1566
 QY 2475 CTTGAGAGGATTTGATGATGATGATGATGATGATGATGATGATGATGATGATGATG 2534
 DB 1567 CAGGCTAGGATTTGATGATGATGATGATGATGATGATGATGATGATGATGATGATG 1626
 QY 2535 ACTGATCTAGGAG 2594
 DB 1627 ACTGATCTAGGAGC---AGATCAGCTTCACTTCACTGAGAGATCACTCGACGCTGT 1683
 QY 2595 GCTGCGCAGATTAAG 2654
 DB 1684 GCTGCGCAG 1743
 QY 2655 GCTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATG 2714
 DB 1744 GCTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATG 1803
 QY 2715 GCTGAGGAGATTAAG 2768
 DB 1804 GCTGAGGAGATTAAG 1857

RESULT 11
 US-10-799-870-24
 ; Sequence 24, Application US/10799870
 ; Publication No. US2004015888A1
 ; GENERAL INFORMATION:
 ; APPLICANT: Wright, David A.
 ; APPLICANT: Voytes, Daniel F.
 ; TITLE OF INVENTION: PLANT RETROBLEMENTS AND METHODS RELATED THERETO

FILE REFERENCE: P-1065A
CURRENT APPLICATION NUMBER: US/10/799,870
CURRENT FILING DATE: 2004-03-12
PRIOR APPLICATION NUMBER: US/09/586,106
PRIOR FILING DATE: 2003-02-07
PRIOR APPLICATION NUMBER: 60/087,125
PRIOR FILING DATE: 1998-05-29
PRIOR APPLICATION NUMBER: 09/322,478
PRIOR FILING DATE: 1999-05-28
NUMBER OF SEQ ID NOS: 190
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 24
LENGTH: 1857
TYPE: DNA
ORGANISM: Arabidopsis thaliana
US-10-799-870-24

Query Match 24.1%; Score 751.6; DB 19; Length 1857;
Best Local Similarity 65.3%; Pred. No. 3.3e-198;
Matches 1250; Conservative 0; Mismatches 564; Indels 100; Gaps 6;

QY 898 ATGAGTAATCACTAGTGAATAATCTCTATGACCCCTGATTATATGATGAGCTAAG 957
DB 1 ATGAGCAATTACAGTGGCAGTTCTTCTGTGATCCTGACTACAACTGATGAGCAGAA 60
QY 958 TCCTGCTCACTAGACCGAGGTAGAGCAACATGTTTACGAGAGCTATAGGATGATTT 1017
DB 61 TCGATCATCTTCAAGCCAGAGAGAGAACAGAGAAATGAAAGTTTCAG----- 110
QY 1018 GAAGCTCTGCAGCTGCAGTATCAAAAGAAAGCTGAAATGCTTAGAGAGAAAGGCG 1077
DB 111 -----AAGAAAGCTGAGATAGCCCGAGAAAGAGAG 144
QY 1078 ATGCGAGTATGATGAGCTGATGATGAGATATCAAAATGAGTATGAGCCAGAGTCA 1137
DB 145 ATGAGAGAGAGTATGAGCTTATAGCAGAGATCTGAGAGACGATGATCTGTAACAG 204
QY 1138 TGGCGCAAGAGACGAAGTACTGAAACAATCCGAGAGTTTACAGTGAAGATATATC 1197
DB 205 ACTCGCAGAGTACCAAACTTCTGCACAAAGCCGACATATCTGCTGAGGATATGTT 264
QY 1198 AGATCTTGAATGATGATCTTGGGGAGACGAGATATCCTGATATGAGCTTTAGC 1257
DB 265 AGGCTTTTCAAGCTAATGATGATCTGAGCAGAGATATCTTGTCTGCACTTCTGCA 324
QY 1258 CAGTTGGGTTACTGAGAGACGTGCAGATCTGTTGAGAGATGTCATCTG----- 1308
DB 325 CAATCGGATTTGTGAGAGATTTGAGCCTGTACCAAAAGTTGTATCTGAGACCTTGG 384
QY 1309 -----ATAGAGAGAGACAAATGAGATTTCTTTCCACACTGCA 1346
DB 385 ATGCGTTATCCGTATGATAGCATATGAAATGAGACAAATCAATTCCTCCACACTACA 444
QY 1347 GTGGAATGATAGAGGAGTCAACAGACTTGGAGTGGATATCAATGGGGTTAGGCTTTG 1406
DB 445 GTAGAGCTCTACCAAGTATGACCTCTGATGATGATTTGATTTGAGGATTTGGATTT 504
QY 1407 AGGTTCTTATGATGAGACAGGGGTAACGATTTTATGATCAAGAAATTTGAGAAAGCTTT 1466
DB 505 CGATTTTCTGTATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 564
QY 1467 GGTTCCTCTAGTGAAGAGGAAACCAACCCAGGTTTGAAGGAGAGGCTTAAAGATTG 1526
DB 565 GATTTTCCAGTGAAGAGGAGCTTAAGCAAAAGTATGAGAAAGAGTGAAGACTTG 624
QY 1527 TGGGCTACTATTTGGGAAACATCTACCGCTAAACTGAGCGCGTCCAAGACCAACATTC 1586
DB 625 TGGATCACCATGAGGAGCTCTGTACCGTTGAATGCTTCCAGATCAAAAGGACATCAGAA 684
QY 1587 CGAGGCTGTGATGCTACTTTGAGGCTGGTGGTGGCAATGTTTAACTCCAGGGAG 1646
DB 685 CGAGGCTGTGATGCTACTTTGAGGCTGGTGGTGGCAATGTTTAACTCCAGGGAG 744

QY 1647 TCTACGGCACCGTGTCTAACACAGACATGAAATGATATGATTCAGCGCTTATAGGATT 1706
DB 745 ATTACAGGAGACTGTACATTAATCTGATATGAGATGATCGAATGCGCTTCAAGAACT 804
QY 1707 CTCGCGCTTCAAAAGGAAAGAAATGCTCTGAGAGAGATCTTAAAGACTCACCAAGTA 1766
DB 805 CTCGCGCAAACTTAAATGGAATGCTCTTCCAGGAGTGAAGTCAATGACACCTCTCT 864
QY 1767 ATGCTCTGTGATCACTGTGTGGGTAATGAAAGTGGGCGCTGACAAACGGAAGAG 1826
DB 865 ATACTTCTTGTATGATCACTGTGTGATACAAAACGCGGCGTCAAGATTAACGCAAG 924
QY 1827 AAGTAAGAGAGACATATGCTGGTGGGCTGTGACGCCAATTTGAAAGTTTGTGA 1886
DB 925 AGAGACAGAGCGCTGTGTGATAGTGGCGTGTGACACTTATTTGATGCTTGTGA 984
QY 1887 GTTCGCTCAAGGAAGTATGAGGTTGACCGGAAATGATGACTTATCACTTGGCGCA 1946
DB 985 GTCCACTCATTTTCTGTGAGCTGAGCTGAGCCAGACAGCAATGATATGACACTTAC 1044
QY 1947 TGTGAGTCTCTGAGTTTGAATGTTGGCGACTTTTCAACGCTACAGGTTGAGCATTTCA 2006
DB 1045 TGCATTTCTGAGATTGCAATGTTGACATTTCCACAGGTTCAAGTTTGAAGACTCT 1104
QY 2007 TCGATTGAATCGCAACATTTCTTTCCCTGCAATTAAGCTACTAGATTTCTGAGGCG 2066
DB 1105 ACAGACAGAGAGCTTACATCTTCTCTGAGCCCTGAGGACACAGGATATGAGGGA 1164
QY 2067 AAGAACTTGACTTCAAGCCCTGAGCTTGAAGATCTTATTTGAGGCGAGTCCGCAACT 2126
DB 1165 GATACATGATTTTATGAGCTGAGATTTGAGCGCTTCTATGAGAAAGCTTCCACATTA 1224
QY 2127 GAGGATTTAGTCAACCGAAGAGCTTACAAATAGAAATGTTGATGACATATGATATA 2186
DB 1225 GATGAGACGATCTTCTTGAAGAGCT-----GTTGAGATGGATGATGAGAT 1275
QY 2187 GATGAGCGGAGTTTGAACAGACATGATATATTTCAAGTGAATATCTTCCAGCGAG 2246
DB 1276 GAGAGATTAAGTTTGAACATGACATGATATATCTTGTGAAACATGATCTTCCAGCGAG 1335
QY 2247 AAAAGCAAGTTTGAAGGAGCTCAACAGAAACAGCAAGCTGACAGAGTGGTCAAG 2306
DB 1336 CAGACAGAGCTTGAATGAGCTTCAATGAAATTAAGTAAATTTGACAGAGTGGTCAAG 1395
QY 2307 AAAAGCAATTAATCTGCAAGTCTTCAAGGCTTCAAGGCTTCAAGTCTTCAAGAGCA 2366
DB 1396 AAGCAGAGACAGCTGATGCAAGG-----TTTCAAGCTTCTGACAGACAGCTG 1446
QY 2367 AGCTGCTCTTCTTCACTACATATTTCCGAAATGACAGCTCCTCAGAGACATGCTTGG 2426
DB 1447 AGTGTCTTCTTCCACACTGTATTTCCACAGGATCAACCTCTATGAAATGCAATCG 1506
QY 2427 AAGGATATGAGCG-----GTCGAGCTGAGAGCAAGATTTCTGATGTC 2474
DB 1507 AAGGAAATTAATGCACTTGCAGAGGCTTGAAGCTTACGAGCAGAGAGTCCACATGTC 1566
QY 2475 CTTGCGAGCAATTCATCTTGAAGCTTGTGATCTGAAATCTGAGAAATGAGAGACGACTC 2534
DB 1567 CAGGCTAGCATTTGTATTTGAAATCCCGGAAACAGAAAGAAAGAGGATCACTC 1626
QY 2535 ACTGATTTAGACAGAGAGACAGACTTTCGAGTCTGTAATTTAGCGAGCGCGT 2594
DB 1627 ACTGATCTTACAGC---AGATACGCTCTCATTTCACTGAGAGAGATCACTGACCGTGT 1683
QY 2595 GCTGCGCAGATTAAGAAAGAGAGTGAAGTATCTTGAAGCGGCTGCGCGCAGAGA 2654
DB 1684 GCTGCGCAGAGAGAGAGAGTGAAGTATCTTGAAGCGGCTGCGCGCAGAGA 1743
QY 2655 GCTGATGATGATGATCAACATGCTGAGCTGATAGAGAAATGCGGCTTGTCTATG 2714
DB 1744 GCTGATGATGATGATCAACATGCTGAGCTGATAGAGAAAGAGGTTGTCTATG 1803
QY 2715 GCTTGGAGCAATCAAGGACGCAATTTGATCACTTCTTATTTTGCAC 2768

Db 613 NMAACCKTYCYWCMYVCKKCAAGWYKNNCTCMACTCTMYNTTCTCTCNKCCCM 672
Qy 302 GACGACCTTCACCATCAACCAATCAATGCTTTCTTCCTTCATTAAGCTTGACATATC 361
Db 673 KACCKNTTCTCWSCCCCCAKAYCAYCMTMTCTCMTCKACSCCCYCNMYCCMM 732
Qy 362 GACGCGTGAACACTTATCACTCAAGCTCTCATCTCATCTGTTTCCACACGCTG 421
Db 733 CMTCTCTMNAKCANCTTCTTCTCTCMYMTMAKCMCNANTONCKSACTTCTCAG 792
Qy 422 CTCTCATCCCCCAAGAACTGTCTCATCTCTCACTCACTCAAGTTCTCATCTCA 481
Db 793 TTKKCKMTCTCTTCKCKYVWCMYCNKCCCTCMTCTCTCTCTCTCTCTCTCTCT 852
Qy 482 GCAACCAACTGACCT 540
Db 853 YAKCKAKMCTCCCAKMKCACTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 912
Qy 541 TCTTCATCT 600
Db 913 TCT 972
Qy 601 CTCCTCATCT 650
Db 973 NYCTCCKAGTYCKCKCKTCKCKYVWCMYCNKCCCTCMTCTCTCTCTCTCTCT 1032
Qy 661 CCTCTCATCT 711
Db 1033 CTCCKMKTCTCCCMCCCATCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 1083

RESULT 15

US-10-437-963-28390
; Sequence 28390, Application US/10437963
; Publication No. US20040123343A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; APPLICANT: Wu, Wei
; APPLICANT: Boukharov, Andrey A.
; APPLICANT: Barbazuk, Brad
; APPLICANT: Li, Ping
; TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53221)B
; CURRENT APPLICATION NUMBER: US/10/437,963
; CURRENT FILING DATE: 2003-05-14
; NUMBER OF SEQ ID NOS: 204966
; SEQ ID NO 28390
; LENGTH: 862
; TYPE: DNA
; ORGANISM: Oryza sativa
; FEATURE:
; NAME/KEY: unsure
; LOCATION: (1)..(862)
; OTHER INFORMATION: unsure at all n locations
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT4530_3295C.1
US-10-437-963-28390

Query Match 2.0%; Score 62; DB 19; Length 862;
Best Local Similarity 44.3%; Pred. No. 5.8e-06;
Matches 239; Conservative 0; Mismatches 300; Indels 0; Gaps 0;

Qy 53 CCGGCATCGCTTCTCTCAAACTCTCACTGACACCGCGCTCTCTCACTTACTCG 112
Db 298 CC 357
Qy 113 CTTCATCGCTCTGCGCATCTCTCAAACTCTCACTGACACCGCGATATCACTGAGCTCG 172

Db 358 CTCCKNCCCCCCCCCCCCCAACCTCTCCCTTTTCCCCCCCCCCCCCCCCCCCCCTCC 417
Qy 173 CCGCTTCTACAGCGCTCTCATCTGATGATGATGATGATGATGATGATGATGATGAT 232
Db 418 TCCCCACCT 477
Qy 233 AGCTTCATTTCTACTCTGATGATGATGATGATGATGATGATGATGATGATGATGAT 292
Db 478 NCCC 537
Qy 293 TAACTGATGACCACTTCACTCACTCACTCACTCACTCACTCACTCACTCACTCA 352
Db 538 CTCCTCCC 597
Qy 353 GACATCTGACCGGTAACATCTATACCTTCAAGCTCTCACTCTCTCTCTCTCTCTCT 412
Db 598 CCGCT 657
Qy 413 ACAAGCT 472
Db 658 CC 717
Qy 473 CTGATTCAGCACTCACTGATGATGATGATGATGATGATGATGATGATGATGATGAT 532
Db 718 CT 777
Qy 533 CCTCAGCATCTTCT 591
Db 778 CT 836

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